SEQUENCE LISTING

<110> Wang, Tongtong
 Marnerakis, Margarita
 Fanger, Gary R.
 Vedvick, Thomas S.
 Carter, Darrick
 Watanabe, Yoshihiro
 Henderson, Robert A.
 Peckham, David W.
 Fanger, Neil

<120> COMPOSITIONS AND METHODS FOR THE THERAPY
AND DIAGNOSIS OF LUNG CANCER

<130> 210121.455C16

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<141> 2001-06-28

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cataactttt aacaacactg ctctgtaatg ggttgaactg tggtactcag actgagataa 180 ctgaaatgag tggatgtata gtgttattgc ataattatcc cactatgaag caaagggact 240 ggataaattc ccagtctaga ttattagcct ttgttaacca tcaagcacct agaagaagaa 300 ttattggaaa ttttgtcctc tgtaactggc actttggggt gtgacttatc ttttgccttt 360

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100
     catcctcacc atacaccatc cactttccaa taacatttaa teetttetaa aattgtaagt 120
     atacaattgt actttctttg gattttcata acaaatatac catagactgt taattttatt 180
4 2 2
     qaaqtttcct taatqqaatq aqtcattttt gtcttgtgct tttgaggtta cctttgcttt 240
gacttccaac aatttgatca tatagtgttg agctgtggaa atctttaagt ttattctata 300
١., أ
     gcaataattt ctattnnnag anncenggnn naaaannann annaaa
                                                                           346
۶.,<u>ڐ</u>
     <210> 4
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12
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171
1
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110
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i zk
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      tetettetee aagttgtget ttgtggggae aatcattett tgaacattag agaggaagge 180
      agttcaagct gttgaaaaga ctattgctta tttttgtttt taaagaccta cttgacgtca 240
      tgtggacagt gcacgtgcct tacgctacat cttgttttct aggaagaagg ggatgcnggg 300
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      aaaacaaaac aa
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gcataaagcc aatgtagtcc agtttctaag atcatgttcc aagctaactg aatcccactt 180
caatacacac tcatgaactc ctgatggaac aataacaggc ccaagcctgt ggtatgatgt 240
qcacacttqc taqactcaqa aaaaatacta ctctcataaa tgggtgggag tattttgggt 300
gacaacctac tttgcttggc tgagtgaagg aatgatattc atatnttcat ttattccatg 360
gacatttagt tagtgetttt tatataccag geatgatget gagtgacaet ettgtgtata 420
tntccaaatn ttngtncngt cgctgcacat atctgaaatc ctatattaag antttcccaa 480
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ctaaaaccnt ctnctnnang gttagacngg acctetette teeetteeeg aanaatnaag 600
tqtqnqaaqa nancenene eeceetnen tnenneetng eengetnnne enentgtngg 660
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717, 723, 724, 725, 733
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catgittatc tittattatg intigigaag tigigtetti teactaatta eetataetat 120
gccaatattt ccttatatct atccataaca tttatactac atttgtaaga gaatatgcac 180
gtgaaactta acactttata aggtaaaaat gaggtttcca agatttaata atctgatcaa 240
qttcttqtta tttccaaata qaatggactt ggtctgttaa ggggctaagg gagaagaaga 300
agataaggtt aaaagttgtt aatgaccaaa cattctaaaa gaaatgcaaa aaaaaattta 360
ttttcaagcc ttcgaactat ttaaggaaag caaaatcatt tcctanatgc atatcatttg 420
tgagantttc tcantaatat cctgaatcat tcatttcagc tnaggcttca tgttgactcg 480
atatqtcatc tagggaaagt ctatttcatg gtccaaacct gttgccatag ttggtnaggc 540
tttcctttaa ntgtgaanta ttnacangaa attttctctt tnanagttct tnatagggtt 600
aggggtgtgg gaaaagcttc taacaatctg tagtgttncg tgttatctgt ncagaaccan 660
aatnacggat cgnangaagg actgggtcta tttacangaa cgaatnatct ngttnnntgt 720
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gtnnncaact ccngggagcc
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<211> 670
<212> DNA
<213> Homo sapiens
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639, 653, 659, 661
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ccaaggtgca ctcggtggcc tggagttgcg acgggcgtcg cctacctcgg ggtcttcgac 240
aagacgccac gtcttcttgc tgganaanga ccgttggtca aagaaaacaa ttatcgggga 300
catggggata gtgtggacca ctttgttggc atccaagtaa tcctgaccta tttgttacgg 360
cgtctggaga taaaaccatt cgcatctggg atgtgaggac tacaaaatgc attgccactg 420
tgaacactaa aggggagaac attaatatct gctggantcc tgatgggcan accattgctg 480
tagenacaag gatgatgtgg tgactttatt gatgecaaga aacceegtte caaageaaaa 540
aaacanttcc aanttcgaag tcaccnaaat ctcctggaac aatgaacatn aatatnttct 600
tectgacaat ggneettggg tgtnteacat ceteagetne eccaaaaetg aaneetgtne 660
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natccacccc
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610, 620, 621, 622, 628, 641, 646, 656, 673
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cacctagcat tgcctactta gccccctgaa ttaacagagc ccaattgaga caaacccctg 180
gcaacaggaa attcaaggga gaaaaagtaa gcaacttggg ctaggatgag ctgactccct 240
tagagcaaag ganagacagc ccccattacc aaataccatt tttgcctggg gcttgtgcag 300
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gatatatntt cctagtggtt tgactttnaa aataaatnag gtttantttt ctccccccnn 480
ennthetnee intenetenn ennteeecce enetengtee teennnnttn gggggggeen 540
cccccncggn ggacccccct ttggtccctt agtggaggtt natggcccct ggnnttatcc 600
nggccntann tttccccqtn nnaaatqntt cccctccca ntcccnccac ctcaanccqq 660
aagcctaagt ttntaccctg ggggtcccc
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<210> 9
<211> 674
<212> DNA
<213> Homo sapiens
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<222> 602, 632, 639, 668
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taaaaaatqc ttqttctata qtqqaqtaaq aqctcacaca cccaaqqcaq caagataact 120
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ataagcctqa aqqqaaqtaq ctatqaqact ttccattttt cttaqttctc ccaataggct 240
ccttcatqqa aaaaqqcttc ctqtaataat tttcacctaa tqaattaqca qtqtqattat 300
ttctgaaata agagacaaat tgggccgcag agtcttcctg tgatttaaaa taaacaaccc 360
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caaaaacatt agctgttctg tctttcaatt tcaagttatt ttggagactg cctccatgtg 480
agttaattac tttgctctgg aactagcatt attgtcatta tcatcacatt ctgtcatcat 540
catctgaata atattgtgga tttccccctc tgcttgcatc ttcttttgac tcctctggga 600
anaaatqtca aaaaaaagg tcgatctact cngcaaggnc catctaatca ctgcgctgga 660
                                                                   674
aggaccenct gccc
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<213> Homo sapiens
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ttctgtctgt aacaaaaatg tactttatag agatggagga aaaggtctaa tactacatag 120
ccttaagtgt ttctgtcatt gttcaagtgt attttctgta acagaaacat atttggaatg 180
tttttctttt ccccttataa attgtaattc ctgaaatact gctgctttaa aaagtcccac 240
tgtcagatta tattatctaa caattgaata ttgtaaatat acttgtctta cctctcaata 300
aaagggtact tttctattan nnagnngnnn gnnnnataaa anaaaa
                                                                   346
<210> 11
<211> 602
<212> DNA
<213> Homo sapiens
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tgcttccctt tatctggaat gtggcattag cttttttatt ttaaccctct ttaattctta 180
ttcaattcca tgacttaagg ttggagagct aaacactggg atttttggat aacagactga 240
cagttttgca taattataat cggcattgta catagaaagg atatggctac cttttgttaa 300
atctgcactt tctaaatatc aaaaaaggga aatgaagtta taaatcaatt tttgtataat 360
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587, 588, 589, 590, 592, 593, 598, 599, 603, 605, 608
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674, 675, 682, 683
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qcatqcattt qtaacatqat taqtaqattt qaatatataq atqtaqtatn ttgggtatct 180
aggtgtttta tcattatgta aaggaattaa agtaaaggac tttgtagttg tttttattaa 240
atatgcatat agtagagtgc aaaaatatag caaaaatana aactaaaggt agaaaagcat 300
tttagatatg ccttaatnta nnaactgtgc caggtggccc tcggaataga tgccaggcag 360
agaccagtgc etgggtggtg ceteceettg tetgeeecee tgaagaaett eeeteaegtg 420
angtagtgcc ctcgtaggtg tcacgtggan tantggganc aggccgnncn gtnanaagaa 480
ancanngtga nagtttenee gtngangeng aactgteect gngeennnae geteecanaa 540
enthteeaat ngacaatega gttteennne teengnaace tngeegnnnn enngeeenne 600
cantnighta accoegogoe eggategete tennniegti etenenenaa ngggnitten 660
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enneegeeqt enenneegeg ennee
<210> 13
<211> 694
<212> DNA
<213> Homo sapiens
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679, 687
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agttgacgaa gatctggttt acaagaacta attaaatgtt tcattgcatt tttgtaagaa 120
cagaataatt ttataaaatg tttgtagttt ataattgccg aaaataattt aaagacactt 180
tttctctqtq tqtqcaaatq tqtqtttqtq atccattttt ttttttttt taggacacct 240
gtttactagc tagctttaca atatgccaaa aaaggatttc teeetgaccc cateegtggt 300
teaccetett tteeceecat getttttgee etagtttata acaaaggaat gatgatgatt 360
taaaaagtag ttctgtatct tcagtatctt ggtcttccag aaccctctgg ttgggaagqq 420
gatcattttt tactggtcat ttccctttgg agtgtactac tttaacagat ggaaagaact 480
cattggccat ggaaacagcc gangtgttgg gagccagcag tgcatggcac cgtccggcat 540
ctggcntgat tggtctggct gccgtcattg tcagcacagt gccatgggac atggggaana 600
ctgactgcac ngccaatggt tttcatgaag aatacngcat ncncngtgat cacgtnancc 660
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<210> 14
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<210> 14

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226, 229, 239, 241, 245, 252, 255, 259, 303, 309, 359, 387,
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592, 609, 610, 618, 620, 626, 627, 633, 639, 645, 654
<223> n = A, T, C or G
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ccaagtgcat caaatacctg engtneggat ntaaattcat ettetggett geegggattg 180
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ccatctgggg atattccact ncgatnatgt gattaaggaa ntccacggag ttttacaagg 480
acacgtacaa cnacctgaaa accnnggatg anccccaccg ggaancnetg aangccatcc 540
actatgcgtt gaactgcaat ggtttggctg gggnccttga acaatttaat cncatacatc 600
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cagaagtctc gaacaatcc
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242, 261, 266, 270, 278, 285, 286, 298, 311, 324, 337, 350,
363, 384, 391, 395, 405, 411, 424, 427, 443, 448, 453, 455,
458, 463, 467, 470, 479, 482, 484, 493, 499, 505, 518
<223> n = A, T, C or G
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<222> 520, 523, 531, 540, 584, 595, 597, 609, 611, 626, 628, 651,
652, 657, 661, 665, 669, 672, 681, 683, 691, 693
<223> n = A, T, C or G
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cattacaact acccaatccg aagtgtcaac tgtgtcagga ctaanaaacc ctggttttga 120
ttaaaaaagg gcctgaaaaa aggggagcca caaatctgtc tgcttcctca cnttantcnt 180
tggcaaatna gcattctgtc tcnttggctg engectcane neaaaaaane ngaactenat 240
enggeecagg aatacatete neaatnaach aaattganea aggenntggg aaatgeenga 300
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conagttotg ttagaaaaat goongaatto naacnooggt tttontacto ngaatttaga 420
tctncanaaa cttcctggcc acnattcnaa ttnanggnca cgnacanatn ccttccatna 480
aneneacece aentttgana geeangacaa tgaetgentn aantgaagge ntgaaggaan 540
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    <222> 299, 354, 483, 555, 571, 573, 577, 642, 651, 662, 667
    <223> n = A, T, C or G
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    ttcccgggcc ccttacactc cacagtcccg gtcccgccat gtcccagaaa caagaagaag 120
    agaaccetge ggaggagace ggegaggaga ageaggacae geaggagaaa gaaggtatte 180
tgcctgagag agctgaagag gcaaagctaa aggccaaata cccaagccta ggacaaaagc 240
    ctggaggctc cgacttcctc atgaagagac tccagaaagg gcaaaagtac tttgactcng 300
gagactacaa catggccaaa gccaacatga agaataagca gctgccaagt gcangaccag 360
1 1 1
    acaagaacct ggtgactggt gatcacatcc ccaccccaca ggatctgccc agagaaagtc 420
1.0
    ctcgctcgtc accagcaagc ttgcgggtgg ccaagttgaa tgatgctgcc ggggctctgc 480
1.4
    canatotgag acgetteect ecetgeecea ecegggteet gtgetggete etgeeettee 540
, J. ...
    tgcttttgca gccangggtc aggaagtggc nenggtngtg gctggaaagc aaaaccttt 600
117
    cctgttggtg tcccacccat ggagcccctg gggcgagccc angaacttga ncctttttgt 660
    tntcttncc
Ξ
105
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113
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100
200
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    141, 143, 150, 156, 166, 167, 170, 172, 180, 181, 190, 192,
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     242, 244, 251, 253, 256, 268, 297, 305, 308, 311, 314
    <223> n = A, T, C or G
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     <222> 315, 317, 322, 324, 327, 333, 337, 343, 362, 364, 367, 368,
     373, 384, 388, 394, 406, 411, 413, 423, 429, 438, 449, 450,
     473, 476, 479, 489, 491, 494, 499, 505, 507, 508, 522, 523,
     527, 530, 533, 535, 538, 539, 545, 548, 550, 552, 555
     <223> n = A, T, C or G
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     628, 632, 638, 642, 644, 653, 658, 662, 663, 665, 669, 675,
     680, 686, 689
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     <400> 17
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geetgeecan ggganeeca neneteggan eccatnteae accegnneen thegeecaen 180
neetggeten enengeeeng neeagetene gneeeeetee geennneten tinnentete 240
enencectee nenacnacet ectaceencg getecetece cageeceece cegeaaneet 300
ccacnaence ntennencga anencenete genetengee cengececet geceeegee 360
enchaenneg egnteeceeg egenegenge eteneceet eccaenaeag nencaecege 420
agneacgene teegecenet gaegeeeenn eeegeegege teacetteat ggneenaeng 480
ccccqctcnc nccnctgcnc gccgncnngg cgccccgccc cnnccgngtn ccncncgnng 540
ccccngcngn angengtgcg enneangnee gngccgnnen neaccetecg neeneegeee 600
cgcccgctgg gggctcccgc cncgcggntc antccccncc cntncgccca ctntccgntc 660
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cnnenetene getengegen egeceneene ecceece
<210> 18
<211> 670
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 234, 292, 329, 437, 458, 478, 487, 524, 542, 549, 550, 557,
576, 597, 603, 604, 646, 665
<223> n = A, T, C or G
<400> 18
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ctgacctcca gtgccgccgg cctcaagatc agacatggcc cagaacttga acgacttggc 120
gggacggctg cccgcgggc cccggggcat gggcacggcc ctgaagctgt tgctgggggc 180
cggcgccgtg gcctacggtg tgcgcgaatc tgtgttcacc gtggaaggcg ggcncagagc 240
catcttcttc aatcggatcg gtggagtgca caggacacta tcctgggccg anggccttca 300
cttcaggatc cttggttcca gtaccccanc atctatgaca ttcgggccag acctcgaaaa 360
aatctcctcc ctacaggctc caaagaccta cagatggtga atatctccct gcgagtgttg 420
tctcgaccaa tgctcangaa cttcctaaca tgttccancg cctaagggct ggactacnaa 480
gaacgantgt tgccgtccat tgtcacgaag tgctcaagaa tttnggtggc caagttcaat 540
gncctcacnn ctgatcnccc agcggggcca agttanccct ggttgatccc cgggganctg 600
acnnaaaagg gccaaggact tcccctcatc ctggataatg tggccntcac aaagctcaac 660
                                                                   670
tttanccacc
<210> 19
 <211> 606
 <212> DNA
 <213> Homo sapiens
 <220>
 <221> misc feature
 <222> 506
 <223> n = A, T, C or G
 <400> 19
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 tgtcgccttg gctcaactgt ggttgatttg tctgtgcccg gaaagtttgg catcattcgt 180
 ccaggetgtg ccctggaaag tactacagee atectecaae agaagtaegg actgeteece 240
 tcacatgcgt cctacctgtg aaactctggg aagcaggaag gcccaagacc tggtgctgga 300
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tactatgtgt ctgtccactg acgactgtca aggcctcatt tgcagaggcc accggagcta 360
gggcactage etgaetttta aggeagtgtg tetttetgag caetgtagae caageeettg 420
gagetgetgg tttageettg cacetgggga aaggatgtat ttatttgtat tttcatatat 480
cagccaaaag ctgaatggaa aagttnagaa cattcctagg tggccttatt ctaataagtt 540
tcttctgtct gttttgtttt tcaattgaaa agttattaaa taacagattt agaatctagt 600
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gagacc
<210> 20
<211> 449
<212> DNA
<213> Homo sapiens
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cagcgccaga gccgaggaga acccccgctc cctgaggagg acctgtccaa actcttcaaa 120
ccaccacage egectgeeag gatggaeteg etgeteattg eaggeeagat aaacaettae 180
tgccagaaca tcaaggagtt cactgcccaa aacttaggca agctcttcat ggcccaggct 240
cttcaagaat acaacaacta agaaaaggaa gtttccagaa aagaagttaa catgaactct 300
tgaagtcaca ccagggcaac tcttggaaga aatatatttg catattgaaa agcacagagg 360
atttctttag tgtcattgcc gattttggct ataacagtgt ctttctagcc ataataaaat 420
aaaacaaaat cttgactgct tgctcaaaa
<210> 21
<211> 409
<212> DNA
<213> Homo sapiens
<400> 21
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tatgttgagt gaaagaacaa acacggagaa catactatgt ggttctcttt atgtaacatt 180
acagaaataa aaacagaggc aaccaccttt gaggcagtat ggagtgagat agactggaaa 240
aaggaaggaa ggaaactcta cgctgatgga aatgtctgtg tcttcattgg gtggtagtta 300
tgtggggata tacatttgtc aaaatttatt gaactatata ctaaagaact ctgcatttta 360
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                                                                 409
<210> 22
<211> 649
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 263, 353, 610, 635, 646
<223> n = A, T, C or G
<400> 22
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tgataaggat ggtacttgca tatggtgaat tactactgtt gacagtttcc gcagaaatcc 120
tatttcagtg gaccaacatt gtggcatggc agcaaatgcc aacattttgt ggaatagcag 180
caaatctaca agagaccctg gttggttttt cgttttgttt tctttgtttt ttcccccttc 240
tcctgaatca gcagggatgg aangagggta gggaagttat gaattactcc ttccagtagt 300
agetetgaag tgteacattt aatateagtt ttttttaaac atgattetag ttnaatgtag 360
aagagagaag aaagaggaag tgttcacttt tttaatacac tgatttagaa atttgatgtc 420
```

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ttatatcagt agttctgagg tattgatagc ttgctttatt tctgccttta cgttgacagt 480
gttgaagcag ggtgaataac taggggcata tatattttt ttttttgtaa gctgtttcat 540
gatgttttct ttggaatttc cggataagtt caggaaaaca tctgcatgtt gttatctagt 600
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ctgaagtten tatecatete attacaacaa aaaeneecag aaeggnttg
<210> 23
<211> 669
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 642, 661
<223> n = A, T, C or G
<400> 23
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tatectetga cageetttgg getgeetegg eeccageage caeageagga ggaggtgaca 180
teacetqteq tqccccctc tqtcaagact ccgacacetg aaccagetga ggtggagact 240
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ctgacacttc tgctgaagtt ggaggacaaa ctgaaccggc acctgagctg tgacctgatg 360
ccaaatgaga atateeeega gttggegget gagetggtge agetgggett cattagtgag 420
gctgaccaga gccggttgac ttctctgcta gaagagactt gaacaagttc aattttgcca 480
ggaacagtac ceteaactea geegetgtea eegteteete ttagagetea etegggeeag 540
gccctgatct gcgctgtggc tgtcctggac gtgctgcacc ctctgtcctt ccccccagtc 600
agtattacct gtgaagccct tccctccttt attattcagg anggctgggg gggctccttg 660
                                                                   669
nttctaacc
<210> 24
<211> 442
<212> DNA
<213> Homo sapiens
<400> 24
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tcactgccat cattaagcat cagtttcaaa attatagcca ttcatgattt actttttcca 120
gatgactate attattetag teetttgaat ttgtaagggg aaaaaaaaca aaaacaaaaa 180
cttacgatgc acttttctcc agcacatcag atttcaaatt gaaaattaaa gacatgctat 240
ggtaatgcac ttgctagtac tacacacttt ggtacaacaa aaaacagagg caagaaacaa 300
cggaaagaga aaagcettee tttgttggee ettaaactga gteaagatet gaaatgtaga 360
gatgatetet gacgatacet gtatgttett attgtgtaaa taaaattget ggtatgaaat 420
                                                                   442
gacctaaaaa aaaaaaaaga aa
<210> 25
<211> 656
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 330, 342, 418, 548, 579, 608
<223> n = A, T, C or G
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<400> 25
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accetaatgg ggeagagat atageeetag eecagtggtg acatgaceae teeetttggg 180
aggectgagg tagaggggag tggtatgtgt tttctcagtg gaagcagcac atgagtgggt 240
gacaggatgt tagataaagg ctctagttag ggtgtcattg tcatttgaga gactgacaca 300
ctcctagcag ctggtaaagg ggtgctggan gccatggagg anctctagaa acattagcat 360
gggctgatct gattacttcc tggcatcccg ctcactttta tgggaagtct tattagangg 420
atgggacagt tttccatatc cttgctgtgg agctctggaa cactctctaa atttccctct 480
attaaaaatc actgccctaa ctacacttcc tccttgaagg aatagaaatg gaactttctc 540
tgacatantt cttggcatgg ggagccagcc acaaatgana atctgaacgt gtccaggttt 600
ctcctganac tcatctacat agaattggtt aaaccctccc ttggaataag gaaaaa
<210> 26
<211> 434
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 395
<223> n = A, T, C or G
<400> 26
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ctaggtgttt ccatctatgt ttcaatctgt ccatctacca ggcctcgcga taaaaacaaa 120
acaaaaaaac gctgccaggt tttagaagca gttctggtct caaaaccatc aggatcctgc 180
caccagggtt cttttgaaat agtaccacat gtaaaaggga atttggcttt cacttcatct 240
aataactgaa ttgtcaggct ttgattgata attgtagaaa taagtagcct tctgttgtgg 300
gaataagtta taatcagtat tcatctcttt gttttttgtc actcttttct ctctaattgt 360
qtcatttgta ctgtttgaaa aatatttctt ctatnaaatt aaactaacct gccttaaaaa 420
aaaaaaaaa aaaa
                                                                   434
<210> 27
<211> 654
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> 505, 533, 563, 592, 613, 635, 638
<223> n = A, T, C or G
<400> 27
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tttatactgc atcetttaca ttagecacta aatacgttat tgettgatga agacetttea 180
cagaatccta tggattgcag catttcactt ggctacttca tacccatgcc ttaaagaggg 240
gcagtttctc aaaagcagaa acatgccgcc agttctcaag ttttcctcct aactccattt 300
quatgtaagg gcagctggcc cccaatgtgg ggaggtccga acattttctg aattcccatt 360
ttcttqttcg cggctaaatg acagtttctg tcattactta gattccgatc tttcccaaag 420
gtgttgattt acaaagaggc cagctaatag cagaaatcat gaccctgaaa gagagatgaa 480
atteaagetg tgageeagge agganeteag tatggeaaag gtettgagaa tengeeattt 540
ggtacaaaaa aaattttaaa gcntttatgt tataccatgg aaccatagaa anggcaaggg 600
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654
aattgttaag aanaatttta agtgtccaga cccanaanga aaaaaaaaaa aaaa
<210> 28
<211> 670
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 101, 226, 274, 330, 385, 392, 397, 402, 452, 473, 476, 532,
534, 538, 550, 583, 595, 604, 613, 622, 643, 669
<223> n = A,T,C or G
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ggaaggggg aaagatatgt gggataaact gagaaaagaa nccaaaaacc tcaacatcca 120
aggcagetta ttegaactet geggeagegg caaeggggeg geggggteee tgeteeegge 180
gttcccggtg ctcctggtgt ctctctcggc agctttagcg acctgncttt ccttctgagc 240
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ttgccaaaat gttaccaatc agtgaccaac cnagcacagc caaaaatcgg acntcngctt 480
tagtccgtct tcacacacag aataagaaaa cggcaaaccc accccacttt tnantttnat 540
tattactaan ttttttctgt tgggcaaaag aatctcagga acngccctgg ggccnccgta 600
ctanagttaa ccnagctagt tncatgaaaa atgatgggct ccncctcaat gggaaagcca 660
                                                                   670
agaaaaagnc
<210> 29
<211> 551
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 336, 474, 504, 511, 522, 523, 524, 540, 547
<223> n = A, T, C \text{ or } G
<400> 29
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ccctctccag ccactgatgg gaaagtattc tccatcagtt ctcaaaaatca gcaagaatct 180
teagtaceag aggtgeetga tgttgeacat ttgeeacttg agaagetggg accetgtete 240
cctcttgact taagtcgtgg ttcagaagtt acagcaccgg tagcctcaga ttcctcttac 300
cqtaatqaat gtcccagggc agaaaaagag gatacncaga tgcttccaaa tccttcttcc 360
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aaaaqtqaaa ttqqqaaqac aaaaqctcaa cagcatttqq taaqqaqaaa aganaaqatq 480
aqqaaqqaaq aqaqaaqaa gacnaagatc nctacggacc gnnncggaag aagaagaagn 540
                                                                    551
aaaaaanaaa a
<210> 30
<211> 684
<212> DNA
<213> Homo sapiens
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<220>
<221> misc feature
<222> 545, 570, 606, 657, 684
<223> n = A, T, C or G
<400> 30
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cgagactcat ttcttggaag catccctggc aaaaatgcag ctgagtacaa ggttatcact 120
gtgatagaac ctggactgct ttttgagata atagagatgc tgcagtctga agagacttcc 180
ageaectete agttgaatga attaatgatg gettetgagt caactttaet ggeteaggaa 240
ccacgagaga tgactgcaga tgtaatcgag cttaaaggga aattcctcat caacttagaa 300
ggtggtgata ttcgtgaaga gtcttcctat aaagtaattg tcatgccgac tacgaaagaa 360
aaatgccccc gttgttggaa gtatacagcg ggagtcttca gatacactgt gtcctcgatg 420
tgcagaagtt gtcagtggga aaatagtatt aacagctcac tcgagcaaga accctcctga 480
cagtactggg ctagaagttt ggatggatta tttacaatat aggaaagaaa gccaagaatt 540
aqqtnatqaq tqqatqaqta aatgqtggan gatggggaat tcaaatcaga attatggaag 600
aagttnttcc tgttactata gaaaggaatt atgtttattt acatgcagaa aatatanatg 660
                                                                    684
tgtggtgtgt accgtggatg gaan
<210> 31
<211> 654
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 326, 582, 651
<223> n = A, T, C \text{ or } G
<400> 31
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aacatettet cagaatgace cagaagttat categtggga getggegtge ttggetetge 120
tttggcagct gtgctttcca gagatggaag aaaggtgaca gtcattgaga gagacttaaa 180
agagectgae agaatagttg gagaatteet geageegggt ggttateatg tteteaaaga 240
ccttggtctt ggagatacag tggaaggtct tgatgcccag gttgtaaatg gttacatgat 300
tcatgatcag ggaaagcaaa tcagangttc agattcctta ccctctgtca gaaaacaatc 360
aagtgcagag tggaagagct ttccatcacg gaagattcat catgagtctc cggaaagcag 420
ctatggcaga gcccaatgca aagtttattg aaggtgttgt gttacagtta ttagaggaag 480
atgatgttgt gatgggagtt cagtacaagg ataaagagac tgggagatat caaggaactc 540
catgetecae tgactgttgt tgeagatggg etttteteea antteaggaa aageetggte 600
tcaataaagt ttctgtatca ctcatttggt tggcttctta tgaagaatgc nccc
<210> 32
<211> 673
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> 376, 545, 627
<223> n = A, T, C or G
<400> 32
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tatcacctga caccaggagt tttcattgga aaaggatttg aacctggtgt tactaacatt 120
ttaaagacca cacaaggaag caaaatcttt ctgaaagaag taaatgatac acttctqqtg 180
aatgaattga aatcaaaaga atctgacatc atgacaacaa atggtgtaat tcatgttgta 240
qataaactcc tctatccaqc agacacacct qttqgaaatq atcaactgct ggaaatactt 300
aataaattaa tcaaatacat ccaaattaag tttgttcgtg gtagcacctt caaagaaatc 360
cccgtgactg tctatnagcc aattattaaa aaatacacca aaatcattga tgggagtgcc 420
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gaaattaaaa gacgcttcag ggagacnccc catgaaggaa ttgccagcca caaaaaaatt 660
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cagggattag aaa
<210> 33
<211> 673
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 325, 419, 452, 532, 538, 542, 571, 600, 616, 651, 653, 672
<223> n = A, T, C or G
<400> 33
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ggatctgttg tttcttttgg gtctcacctc atcagtgtgc atagtggcag aaattataaa 120
qaaqqttqaa aqqaqcaqqq aaaaqatcca gaaqcatgtt agttcgacat catcatcttt 180
tcttgaagta tgatgcatat tgcattattt tatttgcaaa ctaggaattg cagtctgagg 240
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tgaaattatg caactttgat atcatattcc ttgatttaaa ttgggctttt gtgattgant 420
gaaactttat aaagcatatg gtcagttatt tnattaaaaa ggcaaaacct gaaccacctt 480
ctgcacttaa agaagtctaa cagtacaaat acctatctat cttagatgga tntattntt 540
tntattttta aatattgtac tatttatggt nggtggggct ttcttactaa tacacaaatn 600
aatttatcat ttcaanggca ttctatttgg gtttagaagt tgattccaag nantgcatat 660
ttcqctactq tnt
<210> 34
<211> 684
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 414, 472, 480, 490, 503, 507, 508, 513, 523, 574, 575, 598,
659, 662, 675
<223> n = A, T, C or G
<400> 34
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tgatcagggc tggtgtagca tccggttcct ttagtgcagc taactgcatt tgtcactgat 120
gaccaaggag gaaatcacta agacatttga gaagcagtgg tatgaacgtt cttggacaag 180
ccacagttet gageettaac eetgtagttt geacacaaga aegageteea eeteeeette 240
ttcaqqaqqa atctqtqcqq ataqattqqc tqqacttttc aatggttctg ggttgcaagt 300
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tgcctggttg gaaggtacag gtgttcagca ccttcggaaa aagggcataa agtngtqqqq 420
qacaattete agteeaaqaa gaatgeattg accattgetg getatttget tneetagtan 480
gaattggatn catttttgac cangatnntt ctnctatgct ttnttgcaat gaaatcaaat 540
cccqcattat ctacaaqtqq tatgaaqtcc tgcnnccccc agagaggctg ttcaggcnat 600
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cncagaagga atttntttcc tccc
                                                                 684
<210> 35
<211> 614
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
\langle 222 \rangle 17, \overline{2}0, 152, 223, 267, 287, 304, 306, 316, 319, 321, 355,
365, 382, 391, 407, 419, 428, 434, 464, 467, 477, 480, 495,
499, 505, 515, 516, 522, 524, 527, 542, 547, 549, 567, 572,
576, 578
<223> n = A, T, C or G
<400> 35
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teactgeatg aagactgget tgteteagtg tnteaacete accagggetg tetettggte 180
caeacctcgc tecetgttag tgccgtatga cageccccat canatgaect tggccaagte 240
acqqtttctc tgtggtcaat gttggtnggc tgattggtgg aaagtanggt ggaccaaagg 300
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ttccngtttc tcctggccct gngtgggcta nggcctgatt cgggaanatg cctttgcang 420
gaaggganga taantgggat ctaccaattg attctggcaa aacnatntct aagattnttn 480
tgetttatgt ggganacana tetanetete atttnntget gnanatnaca ecetaetegt 540
qntcqancnc qtcttcqatt ttcqqanaca cnccantnaa tactqqcqtt ctqttqttaa 600
                                                                  614
aaaaaaaaa aaaa
<210> 36
<211> 686
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 222, 224, 237, 264, 285, 548, 551, 628, 643, 645, 665, 674
<223> n = A, T, C or G
<400> 36
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gggcggggc ctggagcagc ccgaggcact gcagcagaag ananaaaaga cacgacnaac 240
ctcaqctcqc caqtccqqtc qctnqcttcc cqccqcatqq caatnagaca gacqccqctc 300
acctgctctg ggcacacgcg acccgtggtt gatttggcct tcagtggcat cacccttatg 360
agtatttett aateageget tgeaaagatg gttaacetat getaegeeag ggagataeag 420
qaqactqqat tqqaacattt ttqqqqtcta aaqqtctqtt tqqqgtgcaa cactgaataa 480
ggatgccacc aaagcagcta cagcagctqc agatttcaca qcccaaqtqt qqqatqctqt 540
ctcaqqanat naattqataa cctqqctcat aacacattqt caaqaatqtg gatttcccca 600
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ggatattatt atttgtttac cggggganag gataactgtt tcncntattt taattgaaca 660
aactnaaaca aaanctaagg aaatcc
<210> 37
<211> 681
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 7, 10, 11, 19, 25, 32, 46, 53, 77, 93, 101, 103, 109, 115,
123, 128, 139, 157, 175, 180, 192, 193, 194, 212, 218, 226,
227, 233, 240, 241, 259, 260, 267, 289, 296, 297, 298, 312,
313, 314, 320, 325, 330, 337, 345, 346, 352, 353, 356
<223> n = A, T, C or G
<221> misc feature
<222> 382, 385, 400, 427, 481, 484, 485, 491, 505, 515, 533, 542,
544, 554, 557, 560, 561, 564, 575, 583, 589, 595, 607, 619,
628, 634, 641, 645, 658, 670
<223> n = A, T, C or G
<400> 37
gagacanach naacgtcang agaanaaaag angcatggaa cacaanccag genegatgge 60
cacettecca ecageaneca gegeeecca gengeeecca ngneeggang accangacte 120
cancetgnat caatetgane tetatteetg geocatneet aceteggagg tggangeegn 180
aaaqqtcqca cnnncaqaga aqctqctqcc ancaccancc gccccnnccc tqncgggctn 240
nataggaaac tggtgaccnn gctgcanaat tcatacagga gcacgcgang ggcacnnnct 300
cacactgagt tnnngatgan gcctnaccan ggacctnccc cagcnnattg annacnggac 360
tgcggaggaa ggaagacccc gnacnggatc ctggccggcn tgccaccccc ccacccctag 420
qattatnece ettgaetgag tetetgaggg getaecegaa eeegeeteea tteeetaeca 480
natnntgete nategggaet gaeangetgg ggatnggagg ggetateece eancateeee 540
tnanaccaac agenacngan natngggget eccengggte ggngeaacne teetneacce 600
eggegengge etteggtgnt gteeteente aacnaattee naaanggegg geeeecengt 660
ggactcctcn ttgttccctc c
<210> 38
<211> 687
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 3, 30, 132, 151, 203, 226, 228, 233, 252, 264, 279, 306,
308, 320, 340, 347, 380, 407, 429, 437, 440, 445, 448, 491,
559, 567, 586, 589, 593, 596, 603, 605, 606, 609, 626, 639,
655, 674, 682
<223> n = A, T, C or G
<400> 38
canaaaaaa aaaacatggc cgaaaccagn aagctgcgcg atggcgccac ggcccctctt 60
cteccqqcct qtqtccqqaa qqtttccctc cqaqqcqccc cqqctcccqc aagcqgagga 120
gagggeggga entgeegggg eeggagetea naggeeetgg ggeegetetg eteteeegee 180
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ategeaaggg eggegetaac etnaggeete eeegeaaagg teeeenange ggnggeggeg 240

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gggggctgtg anaaccgcaa aaanaacgct gggcgcgcng cgaacccgtc cacccccgcg 300
aaggananac ttccacagan gcagcgtttc cacagcccan agccacnttt ctagggtgat 360
qcaccccaqt aagttcctqn cqqqqaaqct caccqctqtc aaaaaanctc ttcgctccac 420
cggcgcacna aggggangan ggcangangc tgccgcccgc acaggtcatc tgatcacgtc 480
geoegeceta ntetgetttt gtgaatetee actttgttea acceeaeceg eegttetete 540
ctccttgcgc cttcctctna ccttaanaac cagcttcctc tacccnatng tanttnctct 600
genenngtng aaattaatte ggteeneegg aacetettne etgtggeaac tgetnaaaga 660
                                                                   687
aactgctgtt ctgnttactg cngtccc
<210> 39
<211> 695
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 300, 401, 423, 429, 431, 437, 443, 448, 454, 466, 492, 515,
523, 524, 536, 538, 541, 552, 561, 566, 581, 583, 619, 635,
636, 641, 649, 661, 694
<223> n = A, T, C or G
<400> 39
actagtctgg cctacaatag tgtgattcat gtaggacttc tttcatcaat tcaaaacccc 60
tagaaaaacg tatacagatt atataagtag ggataagatt tctaacattt ctgggctctc 120
tgacccctgc gctagactgt ggaaagggag tattattata gtatacaaca ctgctgttgc 180
cttattagtt ataacatgat aggtgctgaa ttgtgattca caatttaaaa acactgtaat 240
ccaaactttt ttttttaact gtagatcatg catgtgaatg ttaatgttaa tttgttcaan 300
gttgttatgg gtagaaaaaa ccacatgcct taaaatttta aaaagcaggg cccaaactta 360
ttagtttaaa attaggggta tgtttccagt ttgttattaa ntggttatag ctctgtttag 420
aanaaatcna ngaacangat ttngaaantt aagntgacat tattinccag tgactigita 480
atttgaaatc anacacggca cetteegttt tggtnetatt ggnntttgaa tecaanengg 540
ntccaaatct tnttggaaac ngtccnttta acttttttac nanatcttat ttttttattt 600
tggaatggcc ctatttaang ttaaaagggg ggggnnccac naccattent gaataaaact 660
naatatatat ccttggtccc ccaaaattta aggng
                                                                   695
<210> 40
<211> 674
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 403, 428, 432, 507, 530, 543, 580, 583, 591, 604, 608, 621,
624, 626, 639, 672
<223> n = A, T, C or G
<400> 40
actagtagtc agttgggagt ggttgctata ccttgacttc atttatatga atttccactt 60
tattaaataa tagaaaagaa aatcccggtg cttgcagtag agttatagga cattctatgc 120
ttacagaaaa tatagccatg attgaaatca aatagtaaag gctgttctgg ctttttatct 180
tettagetea tettaaataa gtagtaeact tgggatgeag tgegtetgaa gtgetaatea 240
gttgtaacaa tagcacaaat cgaacttagg atgtgtttct tctcttctgt gtttcgattt 300
tgatcaattc tttaattttg ggaacctata atacagtttt cctattcttg gagataaaaa 360
ttaaatggat cactgatatt taagtcattc tgcttctcat ctnaatattc catattctgt 420
```

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attagganaa antacctccc agcacagccc cctctcaaac cccacccaaa accaagcatt 480
tggaatgagt ctcctttatt tccgaantgt ggatggtata acccatatcn ctccaatttc 540
tgnttgggtt gggtattaat ttgaactgtg catgaaaagn ggnaatcttt nctttgggtc 600
aaantttncc ggttaatttg nctngncaaa tccaatttnc tttaagggtg tctttataaa 660
atttgctatt cngg
<210> 41
<211> 657
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 243, 247, 251, 261, 267, 272, 298, 312, 315, 421, 432, 434,
501, 524, 569, 594, 607, 650
<223> n = A, T, C or G
<400> 41
gaaacatgca agtaccacac actgtttgaa ttttgcacaa aaagtgactg tagggatcag 60
gtgatagccc cggaatgtac agtgtcttgg tgcaccaaga tgccttctaa aggctgacat 120
accttgggac cctaatgggg cagagagtat agccctagcc cagtggtgac atgaccactc 180
cctttgggag getgaagtta aagggaatgg tatgtgtttt ctcatggaag cagcacatga 240
atnggtnaca ngatgttaaa ntaaggntct antttgggtg tcttgtcatt tgaaaaantg 300
acacactcct ancanctggt aaaggggtgc tggaagccat ggaagaactc taaaaacatt 360
agcatgggct gatctgatta cttcctggca tcccgctcac ttttatggga agtcttatta 420
naaggatggg ananttttcc atateettge tgttggaact etggaacaet etetaaattt 480
ccctctatta aaaatcactg nccttactac acttectect tganggaata gaaatggace 540
tttctctqac ttagttcttg gcatggganc cagcccaaat taaaatctga cttntccggt 600
ttctccngaa ctcacctact tgaattggta aaacctcctt tggaattagn aaaaacc
<210> 42
<211> 389
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 179, 317, 320
<223> n = A, T, C or G
<400> 42
actagtgctg aggaatgtaa acaagtttgc tgggccttgc gagacttcac caggttgttt 60
cgatagetea cacteetgea etgtgeetgt cacceaggaa tgtettttt aattagaaga 120
caggaagaaa acaaaaacca gactgtgtcc cacaatcaga aacctccgtt gtggcagang 180
ggccttcacc gccaccaggg tgtcccgcca gacagggaga gactccagcc ttctgaggcc 240
atcctgaaga attcctgttt gggggttgtg aaggaaaatc acccggattt aaaaagatgc 300
tgttgcctgc ccgcgtngtn gggaagggac tggtttcctg gtgaatttct taaaagaaaa 360
atattttaag ttaagaaaaa aaaaaaaaa
                                                                   389
<210> 43
<211> 279
<212> DNA
<213> Homo sapiens
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<400> 43
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gtaaaggata aaatgaatga gttctgtcat gattcactat tctagaactt gcatgacctt 120
tactgtgtta gctctttgaa tgttcttgaa attttagact ttctttgtaa acaaataata 180
tgtccttatc attgtataaa agctgttatg tgcaacagtg tggagatcct tgtctgattt 240
aataaaatac ttaaacactg aaaaaaaaaa aaaaaaaaa
<210> 44
<211> 449
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 245, 256, 264, 266, 273, 281, 323, 325, 337, 393
\langle 223 \rangle n = A, T, C or G
<400> 44
actagtagca tcttttctac aacgttaaaa ttgcagaagt agcttatcat taaaaaacaa 60
caacaacaac aataacaata aatectaagt gtaaateagt tattetaece cetaccaagg 120
atatcagcct gttttttccc ttttttctcc tgggaataat tgtgggcttc ttcccaaatt 180
totacageet ettteetett eteatgettg agetteeetg tttgeaegea tgegttgtge 240
aagantgggc tgtttngctt ggantneggt cenagtggaa neatgettte cettgttaet 300
gttggaagaa actcaaacct tcnancccta ggtgttncca ttttgtcaag tcatcactgt 360
atttttgtac tggcattaac aaaaaaagaa atnaaatatt gttccattaa actttaataa 420
                                                                    449
aactttaaaa gggaaaaaaa aaaaaaaaa
<210> 45
<211> 559
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 263
<223> n = A, T, C or G
<400> 45
actagtgtgg gggaatcacg gacacttaaa gtcaatctgc gaaataattc ttttattaca 60
cactcactga agtttttgag tcccagagag ccattctatg tcaaacattc caagtactct 120
ttgagageee ageattaeat caacatgeee gtgeagttea aacegaagte egeaggeaaa 180
tttgaagett tgettgteat teaaacagat gaaggeaaga gtattgetat tegaetaatt 240
ggtgaagctc ttggaaaaaa ttnactagaa tactttttgt gttaagttaa ttacataagt 300
tgtattttgt taactttatc tttctacact acaattatgc ttttgtatat atattttgta 360
tgatggatat ctataattgt agattttgtt tttacaagct aatactgaag actcgactga 420
aatattatgt atctagccca tagtattgta cttaactttt acagggtgaa aaaaaaattc 480
tgtgtttgca ttgattatga tattctgaat aaatatggga atatatttta atgtgggtaa 540
aaaaaaaaa aaaaaggaa
                                                                    559
<210> 46
<211> 731
<212> DNA
<213> Homo sapiens
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<220>
<221> misc feature
<222> 270, 467, 477, 502, 635, 660, 671, 688, 695, 697, 725
<223> n = A, T, C or G
<400> 46
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tcaggttccc taacaattgt ttgaaactga atatatatgt ttatgtatgt gtgtgtgttc 120
actgtcatgt atatggtgta tatgggatgt gtgcagtttt cagttatata tatattcata 180
tatacatatg catatatatg tataatatac atatatacat gcatacactt gtataatata 240
catatatata cacatatatq cacacatatn atcactgagt tccaaagtga gtctttattt 300
ggggcaattg tattetetee etetgtetge teaetgggee tttgcaagae atageaattg 360
cttgatttcc tttggataag agtettatet teggeactet tgaetetage ettaacttta 420
gatttctatt ccaqaatacc tctcatatct atcttaaaac ctaaganggg taaagangtc 480
ataagattgt agtatgaaag antttgctta gttaaattat atctcaggaa actcattcat 540
ctacaaatta aattgtaaaa tgatggtttg ttgtatctga aaaaatgttt agaacaagaa 600
atgtaactgg gtacctgtta tatcaaagaa cctcnattta ttaagtctcc tcatagccan 660
atcettatat ngccctctct gacctgantt aatananact tgaataatga atagttaatt 720
taggnttggg c
                                                                   731
<210> 47
<211> 640
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 5, 28, 106, 153, 158, 173, 176, 182, 189, 205, 210, 214,
225, 226, 229, 237, 260, 263, 269, 277, 281, 282, 322, 337,
338, 354, 365, 428, 441, 443, 456, 467, 476, 484, 503, 508,
554, 567, 575, 579, 588, 601, 606, 609, 611, 621, 636
<223> n = A, T, C or G
<400> 47
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cgttaataac teeteaggte eetgeetgea eagggttttt tettantttg ttgeetaaca 120
gtacaccaaa tgtgacatcc tttcaccaat atngattnct tcataccaca tcntcnatgg 180
anacgactnc aacaattttt tgatnacccn aaanactggg ggctnnaana agtacantct 240
ggagcagcat ggacctgtcn gcnactaang gaacaanagt nntgaacatt tacacaacct 300
ttggtatgtc ttactgaaag anagaaacat gcttctnncc ctagaccacg aggncaaccg 360
caganattgc caatgccaag tccgagcggt tagatcaggt aatacattcc atggatgcat 420
tacatacntt gtccccgaaa nanaagatgc cctaanggct tcttcanact ggtccngaaa 480
acanctacac etggtgettg ganaacanac tetttggaag atcatetgge acaagtteee 540
cccagtgggt tttnccttgg cacctanctt accanatena ttcggaance attetttgcc 600
ntggcnttnt nttgggacca ntcttctcac aactgnaccc
                                                                   640
<210> 48
<211> 257
<212> DNA
<213> Homo sapiens
<400> 48
actagtatat gaaaatgtaa atatcacttg tgtactcaaa caaaagttgg tcttaagctt 60
ccaccttgag cagccttgga aacctaacct gcctctttta gcataatcac attttctaaa 120
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tgattttctt tgttcctgaa aaagtgattt gtattagttt tacatttgtt ttttggaaga 180
ttatatttgt atatgtatca tcataaaata tttaaataaa aagtatcttt agagtgaaaa 240
                                                                   257
aaaaaaaaa aaaaaaa
<210> 49
<211> 652
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 410, 428, 496, 571, 647
<223> n = A, T, C or G
<400> 49
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tccacttatt tgaactctta agtcataaat gtataatgac ttatgaatta gcacagttaa 120
qttgacacta qaaactgccc atttctqtat tacactatca aataggaaac attggaaaga 180
tggggaaaaa aatcttattt taaaatggct tagaaagttt tcagattact ttgaaaattc 240
taaacttctt tctgtttcca aaacttgaaa atatgtagat ggactcatgc attaagactg 300
ttttcaaagc tttcctcaca tttttaaagt gtgattttcc ttttaatata catatttatt 360
ttctttaaag cagctatatc ccaacccatg actttggaga tatacctatn aaaccaatat 420
aacagcangg ttattgaagc agctttctca aatgttgctt cagatgtgca agttgcaaat 480
tttattgtat ttgtanaata caatttttgt tttaaactgt atttcaatct atttctccaa 540
gatgetttte atatagagtg aaatateeea ngataaetge ttetgtgteg tegeatttga 600
cgcataactg cacaaatgaa cagtgtatac ctcttggttg tgcattnacc cc
<210> 50
<211> 650
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 237, 270, 311, 443, 454, 488, 520, 535, 539, 556, 567, 594,
603, 634
<223> n = A, T, C or G
<400> 50
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tgttgagtaa aaaggagatg cccaatattc aaagctgcta aatgttctct ttgccataaa 120
gactccgtgt aactgtgtga acacttggga tttttctcct ctgtcccgag gtcgtcgtct 180
gctttctttt ttgggttctt tctagaagat tgagaaatgc atatgacagg ctgagancac 240
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ggctcctgga nggctgcctg ggggaggcag acatgggagt gccaaggtgg ccagatggtt 360
ccaggactac aatgtettta tttttaactg tttgccactg etgeecteae eeetgeeegg 420
ctctggagta ccgtctgccc canacaagtg ggantgaaat gggggtgggg gggaacactg 480
attcccantt agggggtgcc taactgaaca gtagggatan aaggtgtgaa cctgngaant 540
gcttttataa attatnttcc ttgttanatt tattttttaa tttaatctct gttnaactgc 600
ccngggaaaa ggggaaaaaa aaaaaaaaat tctntttaaa cacatgaaca
<210> 51
<211> 545
<212> DNA
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<213> Homo sapiens
<220>
<221> misc feature
<222> 66, 159, 195, 205, 214, 243, 278, 298, 306, 337, 366, 375,
382, 405, 446, 477, 492, 495, 503, 507, 508, 521, 537
<223> n = A,T,C or G
<400> 51
tgqcgtgcaa ccagggtagc tgaaqtttgg gtctgggact ggagattggc cattaggcct 60
cetganatte cagetecett ceaceaagee cagtettget aegtggeaca gggeaaacet 120
gactocottt gggootcagt ttocootooc ottoatgana tgaaaagaat actacttttt 180
cttgttggtc taacnttgct ggacncaaag tgtngtcatt attgttgtat tgggtgatgt 240
qtncaaaact qcaqaaqctc actqcctatq aqaqqaanta aqaqaqataq tqqatqanaq 300
ggacanaagg agtcattatt tggtatagat ccaccentee caacetttet etecteagte 360
cctgcncctc atgtntctgg tntggtgagt cctttgtgcc accanccatc atgctttgca 420
ttgctgccat cctgggaagg gggtgnatcg tctcacaact tgttgtcatc gtttganatg 480
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caaaa
                                                                    545
<210> 52
<211> 678
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
\langle 222 \rangle 98, \overline{1}19, 121, 131, 136, 139, 140, 142, 143, 163, 168, 172,
176, 184, 189, 190, 191, 200, 201, 205, 207, 221, 223, 229,
230, 237, 240, 241, 255, 264, 266, 267, 276, 280, 288, 289,
291, 297, 301, 306, 308, 314, 315, 326, 332, 335, 337
<223> n = A, T, C or G
<221> misc feature
<222> 339, 341, 343, 344, 345, 347, 350, 355, 356, 358, 362, 363,
372, 379, 395, 397, 398, 400, 403, 412, 414, 421, 423, 431,
435, 438, 439, 450, 457, 463, 467, 471, 474, 480, 483, 484,
487, 490, 491, 492, 493, 499, 500, 504, 508, 518, 536
<223> n = A, T, C or G
<221> misc feature
<222> 538, 549, 551, 552, 554, 556, 557, 562, 563, 567, 571, 572,
576, 579, 590, 592, 595, 598, 606, 609, 613, 620, 622, 624,
626, 631, 634, 638, 641, 647, 654, 660, 661, 674
<223> n = A, T, C or G
<400> 52
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ggaggaagac gatttggggg gggagggggg gggggcangg tccgtggggc tttccctant 120
ntateteeat ntecantgnn enntgtegee tetteeeteg teneattnga anttanteee 180
tggnccccnn necetetccn nectnenect ecceetceg nenectecnn etttttntan 240
nettececat eteenteece cetnanngte ceaacneegn cageaatnne neaettnete 300
nctcenence technologit ettethttet enaenththe nennntheen tgeennthaa 360
annotetece energeaane gattetetee eteenennan ethteeaete enthettete 420
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nenegeteet nttentenne ceaecteten cettegnece cantaenete neenecettn 480
egnntentth nunteetenn acenecenee teeettenee eetettetee eeggththte 540
tetetecene nnenenneet ennecentee nngegneent tteegeeeen enceneentt 600
cettentene cantecaten entntnecat netnectnee neteacneee getneeeeen 660
ntctctttca cacnqtcc
<210> 53
<211> 502
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 139, 146, 215, 217, 257, 263, 289, 386, 420, 452, 457, 461,
466, 482, 486
<223> n = A, T, C \text{ or } G
<400> 53
tgaagateet ggtgtegeea tgggeegeeg eeeegeeegt tgttaeeggt attgtaagaa 60
caageegtae ccaaagtete gettetgeeg aggtgteeet gatgeeaaaa ttegeatttt 120
tgacctgggg cggaaaaang caaaantgga tgagtctccg ctttgtggcc acatggtgtc 180
agatcaatat gagcagctgt cctctgaagc cctgnangct gcccgaattt gtgccaataa 240
gtacatggta aaaagtngtg gcnaagatgc ttccatatcc gggtgcggnt ccaccccttc 300
cacgteatee geateaacaa gatgttgtee tgtgetgggg etgacagget eecaacagge 360
atgcgaagtg cetttggaaa acceanggea etgtggeeag ggtteacatt gggeeaattn 420
atcatgttca tccgcaccaa ctgcagaaca angaacntgt naattnaagc cctgcccagg 480
gncaanttca aatttcccgg cc
                                                                   502
<210> 54
<211> 494
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 431, 442, 445
<223> n = A,T,C or G
<400> 54
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tttaatgeca aaagtttget ttgteeacaa ttteettaag acetetteag aaagggattt 120
gtttgcctta atgaatactg ttgggaaaaa acacagtata atgagtgaaa agggcagaag 180
caagaaattt ctacatctta gcgactccaa gaagaatgag tatccacatt tagatggcac 240
attatgagga etttaatett teettaaaca caataatgtt ttettttte ttttatteac 300
atgatttcta agtatatttt tcatgcagga cagtttttca accttgatgt acagtgactg 360
tgttaaattt ttctttcagt ggcaacctct ataatcttta aaatatggtg agcatcttgt 420
ctgttttgaa ngggatatga cnatnaatct atcagatggg aaatcctgtt tccaagttag 480
aaaaaaaaa aaaa
                                                                   494
<210> 55
<211> 606
<212> DNA
<213> Homo sapiens
```

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<220>
<221> misc feature
<222> 375, 395, 511, 542, 559, 569, 578, 581
<223> n = A,T,C or G
<400> 55
actagtaaaa agcagcattg ccaaataatc cctaattttc cactaaaaat ataatgaaat 60
gatgttaagc tttttgaaaa gtttaggtta aacctactgt tgttagatta atgtatttgt 120
tgcttccctt tatctggaat gtggcattag cttttttatt ttaaccctct ttaattctta 180
ttcaattcca tgacttaagg ttggagagct aaacactggg atttttggat aacagactga 240
caqttttqca taattataat cggcattqta cataqaaaqq atatqqctac cttttqttaa 300
atotgoactt totaaatato aaaaaaggga aatgaagtat aaatcaattt ttgtataato 360
tgtttgaaac atgantttta tttgcttaat attanggctt tgcccttttc tgttagtctc 420
ttgggateet gtgtaaaact gtteteatta aacaccaaac agttaagtee attetetggt 480
actagetaca aatteegttt catattetae ntaacaattt aaattaactg aaatatttet 540
anatggtcta cttctgtcnt ataaaaacna aacttgantt nccaaaaaaa aaaaaaaaa 600
aaaaaa
                                                                   606
<210> 56
<211> 183
<212> DNA
<213> Homo sapiens
<400> 56
actagtatat ttaaacttac aggettattt gtaatgtaaa ccaccatttt aatgtactgt 60
aattaacatg gttataatac gtacaatcct tccctcatcc catcacacaa ctttttttgt 120
gtgtgataaa ctgattttgg tttgcaataa aaccttgaaa aataaaaaaaa aaaaaaaaa 180
aaa
                                                                   183
<210> 57
<211> 622
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 358, 368, 412, 414, 425, 430, 453, 455, 469, 475, 495, 499,
529, 540, 564, 575, 590
<223> n = A, T, C or G
<400> 57
actagteact actgtettet cettgtaget aateaateaa tattetteee ttgeetgtgg 60
gcagtggaga gtgctgctgg gtgtacgctg cacctgccca ctgagttggg gaaagaggat 120
aatcagtgag cactgttctg ctcagagctc ctgatctacc ccacccccta ggatccagga 180
ctgggtcaaa gctgcatgaa accaggccct ggcagcaacc tgggaatggc tggaggtggg 240
agagaacctg acttetettt eceteteeet ecteeaacat taetggaact etateetgtt 300
agggatette tgagettgtt teeetgetgg gtgggacaga agacaaagga gaagggangg 360
tctacaanaa gcagcccttc tttgtcctct ggggttaatg agcttgacct ananttcatg 420
gaganaccan aagcetetga tttttaattt eentnaaatg tttgaagtnt atatntacat 480
atatatattt ctttnaatnt ttgagtcttt gatatgtctt aaaatccant ccctctgccn 540
qaaacctqaa ttaaaaccat qaanaaaaat qtttncctta aaqatqttan taattaattq 600
aaacttgaaa aaaaaaaaaa aa
                                                                   622
```

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<211> 433
<212> DNA
<213> Homo sapiens
<400> 58
gaacaaattc tgattggtta tgtaccgtca aaagacttga agaaatttca tgattttgca 60
gtgtggaagc gttgaaaatt gaaagttact gcttttccac ttgctcatat agtaaaggga 120
teettteage tgeeagtgtt gaataatgta teateeagag tgatgttate tgtgaeagte 180
accagettta agetgaacca ttttatgaat accaaataaa tagaeetett gtaetgaaaa 240
catatttgtg actttaatcg tgctgcttgg atagaaatat ttttactggt tcttctgaat 300
tgacagtaaa cetgtecatt atgaatggee tactgtteta ttatttgttt tgacttgaat 360
ttatccacca aagacttcat ttgtgtatca tcaataaagt tgtatgtttc aactgaaaaa 420
aaaaaaaaa aaa
                                                                   433
<210> 59
<211> 649
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 22, 190, 217, 430, 433, 484, 544, 550, 577, 583, 594
<223> n = A, T, C or G
<400> 59
actagttatt atctgacttt enggttataa teattetaat gagtgtgaag tageetetgg 60
tgtcatttgg atttgcattt ctctgatgag tgatgctatc aagcaccttt gctggtgctg 120
ttggccatat gtgtatgttc cctggagaag tgtctgtgct gagccttggc ccacttttta 180
attaggegtn tgtettttta ttactgagtt gtaagantte tttatatatt etggatteta 240
gaccettate agatacatgg tttgcaaata tttteteeca ttetgtgggt tgtgttttea 300
ctttatcgat aatgtcctta gacatataat aaatttgtat tttaaaagtg acttgatttg 360
ggctgtgcaa ggtgggctca cgcttgtaat cccagcactt tgggagactg aggtgggtgg 420
atcatatgan gangctagga gttcgaggtc agcctggcca gcatagcgaa aacttgtctc 480
tacnaaaaat acaaaaatta gtcaggcatg gtggtgcacg tctgtaatac cagcttctca 540
ggangctgan gcacaaggat cacttgaacc ccagaangaa gangttgcag tganctgaag 600
atcatgccag ggcaacaaaa atgagaactt gtttaaaaaa aaaaaaaaa
                                                                   649
<210> 60
<211> 423
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 209, 222, 277, 389, 398
<223> n = A, T, C or G
<400> 60
actagttcag gccttccagt tcactgacaa acatggggaa gtgtgcccag ctggctggaa 60
acctggcagt gataccatca agcctgatgt ccaaaagagc aaagaatatt tctccaagca 120
gaagtgageg etgggetgtt ttagtgeeag getgeggtgg geageeatga gaacaaaace 180
tettetgtat tttttttte cattagtana acacaagaet engatteage egaattgtgg 240
tgtcttacaa ggcagggctt tcctacaggg ggtgganaaa acagcctttc ttcctttggt 300
aggaatggcc tgagttggcg ttgtgggcag gctactggtt tgtatgatgt attagtagag 360
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caacccatta atcttttgta gtttgtatna aacttganct gagaccttaa acaaaaaaaa 420
                                                                423
aaa
<210> 61
<211> 423
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 195, 285, 295, 329, 335, 340, 347, 367, 382, 383, 391, 396,
418
<223> n = A, T, C or G
<400> 61
egggaetgga atgtaaagtg aagtteggag etetgageae gggetettee egeegggtee 60
caggitetgag tatggetggg agteggggge cacaggeete tagetgtget geteaagaag 180
actggatcag ggtanctaca agtggccggg ccttgccttt gggattctac cctgttccta 240
atttggtgtt ggggtgcggg gtccctggcc cccttttcca cactnectcc ctccngacag 300
caacctccct tggggcaatt gggcctggnt ctccncccgn tgttgcnacc ctttgttggt 360
ttaaggnett taaaaatgtt anntttteee ntgeengggt taaaaaagga aaaaactnaa 420
                                                                423
aaa
<210> 62
<211> 683
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 218, 291, 305, 411, 416, 441, 443, 453, 522, 523, 536, 542,
547, 566, 588, 592, 595, 603, 621, 628, 630, 632, 644, 645,
648, 655, 660, 672, 674, 676, 677, 683
<223> n = A,T,C or G
<400> 62
gctggagagg ggtacggact ttcttggagt tgtcccaggt tggaatgaga ctgaactcaa 60
gaagagaccc taagagactg gggaatggtt cctgccttca ggaaagtgaa agacgcttag 120
getgteaaca ettaaaggaa gteeeettga ageeeagagt ggaeagaeta gaeeeattga 180
tggggccact ggccatggtc cgtggacaag acattcengt gggccatggc acaccggggg 240
ggatcaaaat gtgtacttgt ggggtetege eeettgeeaa aaccaaacca nteceactee 300
tgtenttgga etttetteee atteeeteet eeceaaatge aetteeeete eteeetetge 360
coctcotgtg tttttggaat totgtttooc toaaaattgt taatttttta nttttngaco 420
atgaacttat gtttggggtc nangttcccc ttnccaatgc atactaatat attaatggtt 480
atttattttt qaaatatttt ttaatqaact tqqaaaaaat tnntqqaatt tccttncttc 540
cnttttnttt gggggggtg gggggntggg ttaaaatttt tttggaancc cnatnggaaa 600
ttnttacttg gggccccct naaaaaantn anttccaatt cttnnatngc ccctnttccn 660
                                                                 683
ctaaaaaaaa ananannaaa aan
<210> 63
<211> 731
<212> DNA
<213> Homo sapiens
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<220>
<221> misc feature
<222> 237, 249, 263, 288, 312, 317, 323, 326, 337, 352, 362, 370,
377, 400, 411, 414, 434, 436, 446, 457, 473, 486, 497, 498,
502, 512, 531, 546, 554, 563, 565, 566, 588, 597, 608, 611,
613, 615, 627, 632, 640, 641, 644, 654, 660, 663, 665
<223> n = A, T, C or G
<221> misc feature
<222> 671, 678, 692, 697, 698, 699, 704, 705, 712, 714, 717, 718,
719, 723, 725, 730, 731
<223> n = A, T, C or G
<400> 63
actagtcata aagggtgtgc gcgtcttcga cgtggcggtc ttggcgccac tgctgcgaga 60
eccggeeetg gaeeteaagg teateeactt ggtgegtgat eeccgegegg tggegagtte 120
acggateege tegegeeaeg geeteateeg tgagageeta caggtggtge geageegaga 180
ccgcgagctc accgcatgcc cttcttggag gccgcgggcc acaagcttgg cgcccanaaa 240
gaaggegtng ggggeeegea aantaceaeg etetgggege tatggaangt eetettgeaa 300
taatattggt tnaaaanctg canaanagcc cctgcanccc cctgaactgg gntgcagggc 360
cncttacctn gtttggntgc ggttacaaag aacctgtttn ggaaaaccct nccnaaaacc 420
ttccgggaaa attntncaaa tttttnttgg ggaattnttg ggtaaacccc ccnaaaatgg 480
gaaacntttt tgccctnnaa antaaaccat tnggttccgg gggcccccc ncaaaaccct 540
tttttntttt tttntgcccc cantnncccc ccggggcccc tttttttngg ggaaaanccc 600
coccetnce nanantttta aaaqqqnqqq anaatttttn nttnccccc qqqncccccn 660
ggngntaaaa nggtttcncc cccccgaggg gnggggnnnc ctcnnaaacc cntntcnnna 720
concuttttn n
                                                                   731
<210> 64
<211> 313
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 240
<223> n = A, T, C or G
<400> 64
actagttgtg caaaccacga ctgaagaaag acgaaaagtg ggaaataact tgcaacgtct 60
gttagagatg gttgctacac atgttgggtc tgtagagaaa catcttgagg agcagattgc 120
taaagttgat agagaatatg aagaatgcat gtcagaagat ctctcggaaa atattaaaga 180
gattagagat aagtatgaga agaaagctac tctaattaag tcttctgaag aatgaagatn 240
aaatgttgat catgtatata tatccatagt gaataaaatt gtctcagtaa agttgtaaaa 300
                                                                   313
aaaaaaaaa aaa
<210> 65
<211> 420
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
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<222> 400, 402, 403, 404, 405, 406, 409, 411, 412, 414, 415, 416
<223> n = A, T, C or G
<400> 65
actagttccc tggcaggcaa gggcttccaa ctgaggcagt gcatgtgtgg cagagagagg 60
caggaagetg geagtggeag ettetgtgte tagggagggg tgtggeteee teetteeetg 120
totgggaggt tggagggaag aatctaggco ttagcttgco ctcctgccac cottccctt 180
gtagatactg ccttaacact ccctcctct tcagctgtgg ctgccaccca agccaggttt 240
ctccgtgctc actaatttat ttccaggaaa ggtgtgtgga agacatgagc cgtgtataat 300
atttgtttta acattttcat tgcaagtatt gaccatcatc cttggttgtg tatcgttgta 360
acacaaatta atqatattaa aaagcatcca aacaaagccn annnnnaana nnannnqaaa 420
<210> 66
<211> 676
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 328, 454, 505, 555, 586, 612, 636, 641
<223> n = A, T, C or G
<400> 66
actagtttcc tatgatcatt aaactcattc tcagggttaa gaaaggaatg taaatttctg 60
cctcaatttg tacttcatca ataagttttt gaagagtgca gatttttagt caggtcttaa 120
aaataaacte acaaatetgg atgeatttet aaattetgea aatgttteet ggggtgaett 180
aacaaggaat aatcccacaa tatacctagc tacctaatac atggagctgg ggctcaaccc 240
actgttttta aggatttgcg cttacttgtg gctgaggaaa aataagtagt tccgagggaa 300
gtagttttta aatgtgagct tatagatngg aaacagaata tcaacttaat tatggaaatt 360
gttagaaacc tgttctcttg ttatctgaat cttgattgca attactattg tactggatag 420
actocagooo attgcaaagt ctcagatato ttanctgtgt agttgaatto cttggaaatt 480
ctttttaaga aaaaattgga gtttnaaaga aataaacccc tttgttaaat gaagcttggc 540
tttttggtga aaaanaatca tcccgcaggg cttattgttt aaaaanggaa ttttaagcct 600
ccctggaaaa anttgttaat taaatgggga aaatgntggg naaaaattat ccgttagggt 660
ttaaagggaa aactta
                                                                   676
<210> 67
<211> 620
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 419, 493, 519, 568, 605, 610
<223> n = A, T, C or G
<400> 67
caccattaaa getgettaee aagaaettee ceageatttt gaetteettg tttgataget 60
gaattgtgag caggtgatag aagagcettt etagttgaac atacagataa tttgetgaat 120
acattccatt taatqaaqqq qttacatctq ttacqaaqct actaaqaaqq aqcaaqaqca 180
taggggaaaa aaatctgatc agaacgcatc aaactcacat gtgccccctc tactacaaac 240
agattgtagt gctgtggtgg tttattccgt tgtgcagaac ttgcaagctg agtcactaaa 300
cccaaagaga ggaaattata ggttagttaa acattgtaat cccaggaact aagtttaatt 360
```

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cacttttgaa gtgttttgtt ttttattttt ggtttgtctg atttactttg ggggaaaang 420
ctaaaaaaaa agggatatca atctctaatt cagtgcccac taaaagttgt ccctaaaaag 480
tetttaetgg aanttatggg actttttaag etceaggtnt tttggteete caaattaace 540
ttgcatgggc cccttaaaat tgttgaangg cattcctgcc tctaagtttg gggaaaattc 600
ccccnttttn aaaatttgga
                                                                620
<210> 68
<211> 551
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 286, 464, 480, 501, 502, 518, 528, 533, 536, 537, 538, 539,
540, 541, 543, 544, 545, 547, 548, 549
<223> n = A, T, C or G
<400> 68
actagtaget ggtacataat cactgaggag ctatttetta acatgetttt atagaccatg 60
ctaatqctag accagtattt aagggctaat ctcacacctc cttagctgta agagtctggc 120
ttagaacaga cetetetgtg caataacttg tggccactgg aaatecetgg geeggcattt 180
gtattggggt tgcaatgact cccaagggcc aaaagagtta aaggcacgac tgggatttct 240
tetgagaetg tggtgaaaet eetteeaagg etgaggggt eagtangtge tetgggaggg 300
actoggoacc actttgatat toaacaagoc acttgaagoc caattataaa attgttattt 360
tacagetgat ggaactcaat ttgaacette aaaactttgt tagtttatee tattatattg 420
ttaaacctaa ttacatttqt ctaqcattqq atttqqttcc tqtnqcatat qtttttttcn 480
cctatgtgct cccctcccc nnatcttaat ttaaaccnca attttgcnat tcnccnnnnn 540
nannnannna a
                                                                551
<210> 69
<211> 396
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 235, 310, 323, 381
<223> n = A, T, C or G
<400> 69
cagaaatgga aagcagagtt ttcatttctg tttataaacg tctccaaaca aaaatggaaa 60
gcagagtttt cattaaatcc ttttaccttt tttttttctt ggtaatcccc tcaaataaca 120
aattaagcaa atgttaaaag ttttatatgc tttattaatg ttttcaaaag gtatnataca 240
tgtgatacat titttaaget teagitgett giettetggt actitetgit atgggetitt 300
ggggagccan aaaccaatct acnatctett tttgtttgcc aggacatgca ataaaattta 360
                                                                 396
aaaaaataaat aaaaactatt nagaaattga aaaaaa
<210> 70
<211> 536
<212> DNA
<213> Homo sapiens
<220>
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<211> 560

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<221> misc feature
<222> 388, 446, 455
<223> n = A, T, C or G
<400> 70
actagtgcaa aagcaaatat aaacatcgaa aaggcgttcc tcacgttagc tgaagatatc 60
cttcqaaaqa cccctgtaaa aqaqcccaac agtgaaaatg tagatatcag cagtggagga 120
ggcgtgacag gctggaagag caaatgctgc tgagcattct cctgttccat cagttgccat 180
ccactacccc qttttctctt cttqctqcaa aataaaccac tctqtccatt tttaactcta 240
aacagatatt tttgtttctc atcttaacta tccaagccac ctattttatt tgttctttca 300
tetgtgaetg ettgetgaet ttateataat tttetteaaa caaaaaaatg tatagaaaaa 360
tcatgtctgt gacttcattt ttaaatgnta cttgctcagc tcaactgcat ttcagttgtt 420
ttatagtcca gttcttatca acattnaaac ctatngcaat catttcaaat ctattctgca 480
aattgtataa gaataaaagt tagaatttaa caattaaaaa aaaaaaaaa aaaaaa
<210> 71
<211> 865
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 22, 35, 39, 56, 131, 138, 146, 183, 194, 197, 238, 269, 277,
282, 297, 316, 331, 336, 340, 341, 346, 349, 370, 376, 381,
382, 392, 396, 397, 401, 433, 444, 445, 454, 455, 469, 472,
477, 480, 482, 489, 497, 499, 511, 522, 526, 527
<223> n = A, T, C or G
<221> misc feature
<222> 545, 553, 556, 567, 574, 580, 610, 613, 634, 638, 639, 663,
672, 689, 693, 694, 701, 704, 713, 723, 729, 732, 743, 744,
749, 761, 765, 767, 769, 772, 774, 780, 783, 788, 792, 803,
810, 824, 840, 848
<223> n = A, T, C or G
<400> 71
gacaaagegt taggagaaga anagaggeag ggaanactne ceaggeaega tggeeneett 60
cccaccagca accagegece eccaccagee eccaggeeg gaegaegaag actecateet 120
ggattaatet nacetetnte geetgneeca tteetaeete ggaggtggag geeggaaagg 180
tencaccaag aganaanetg etgecaacae caacegeece ageeetggeg ggeaeganag 240
gaaactggtg accaatctgc agaattctna gaggaanaag cnaggggccc cgcgctnaga 300
cagagetgga tatgangcca gaccatggac nctacnecen ncaatneana egggaetgeg 360
gaagatggan gaccenegae nngateagge engetnneea neecceeace ectatgaatt 420
attecegetg aangaatete tgannggett eeannaaage geeteeeene enaaegnaan 480
tncaacatng ggattanang ctgggaactg naaggggcaa ancetnnaat atccccagaa 540
acaanctoto conaanaaac tggggcnoot catnggtggn accaactatt aactaaaccg 600
cacqccaagn aantataaaa ggggggcccc tccncggnng accccctttt gtcccttaat 660
ganggttate encettgegt accatggtne cennttetgt ntgnatgttt ceneteceet 720
concetatnt enageegaac tennatttne eegggggtge natenantng thencetttn 780
ttngttgncc engecettte egneggaach egttteeeeg ttantaaegg eacceggggn 840
aagggtgntt ggcccctcc ctccc
<210> 72
```

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<212> DNA
     <213> Homo sapiens
     <220>
     <221> misc feature
     <222> 83, 173, 183, 186, 209, 211, 215, 255, 321, 322, 323, 335,
     344, 357, 361, 368, 394, 412, 415, 442, 455, 469, 472, 475,
     487, 513, 522, 528, 531, 534, 546
     <223> n = A, T, C or G
     <400> 72
     cetqqacttq tettqqttee aqaacetqae qaceeqqeqa eqqeqaeqte tettttqaet 60
     aaaagacagt gtccagtgct congectagg agtctacggg gaccgcctcc cgcgccgcca 120
     ccatgcccaa cttctctggc aactggaaaa tcatccgatc ggaaaacttc gangaattgc 180
     tenaantget gggggtgaat gtgatgetna ngaanattge tgtggetgea gegteeaage 240
     cagcagtgga gatchaacag gagggagaca ctttctacat caaaacctcc accaccgtgc 300
     gcaccacaaa gattaacttc nnngttgggg aggantttga ggancaaact gtggatngga 360
     ngcctgtnaa aacctggtga aatgggagaa tganaataaa atggtctgtg ancanaaact 420
124
111
     cctgaaagga gaaggccccc anaactcctg gaccngaaaa actgacccnc cnatngggga 480
actgatnett gaaccetgaa egggegggat ganeettttt thttgeenee naangggtte 540
tttccntttc cccaaaaaaa
1,3
     <210> 73
1.4
     <211> 379
.
پرياد
     <212> DNA
<213> Homo sapiens
3 2 M
     <220>
100
     <221> misc feature
     <222> 8, 17, 18, 21, 26, 29, 30, 32, 53, 56, 67, 71, 81, 102, 104,
113
     111, 112, 114, 119, 122, 124, 125, 134, 144, 146, 189, 190,
113
     214, 215, 219, 220, 235, 237, 246, 280, 288, 302, 310, 313,
1 24
     319, 322, 343, 353, 354
į "â
     <223> n = A, T, C or G
     <400> 73
     ctggggancc ggcggtnngc nccatntcnn gncgcgaagg tggcaataaa aancenctga 60
     aaccgcncaa naaacatgcc naagatatgg acgaggaaga tngngctttc nngnacaanc 120
     gnanngagga acanaacaaa ctcnangagc tctcaagcta atgccgcggg gaaggggccc 180
     ttggccacnn gtggaattaa gaaatctggc aaanngtann tgttccttgt gcctnangag 240
     ataagngacc ctttatttca tctgtattta aacctctctn ttccctgnca taacttcttt 300
     tnccacgtan agntggaant anttgttgtc ttggactgtt gtncatttta gannaaactt 360
                                                                          379
     ttgttcaaaa aaaaaataa
     <210> 74
     <211> 437
     <212> DNA
     <213> Homo sapiens
     <220>
     <221> misc feature
     <222> 145, 355
     <223> n = A, T, C or G
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<400> 74
actaqttcaq actqccacqc caaccccaqa aaatacccca catqccaqaa aaqtqaaqtc 60
ctaggtgttt ccatctatgt ttcaatctgt ccatctacca ggcctcgcga taaaaacaaa 120
acaaaaaaac getgecaggt tttanaagca gttetggtet caaaaccate aggateetge 180
caccagggtt cttttgaaat agtaccacat gtaaaaggga atttggcttt cacttcatct 240
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aaaaaaaaa aaaaaaa
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<212> DNA
<213> Homo sapiens
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<222> 440, 513, 539, 551
<223> n = A, T, C or G
<400> 75
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ccctgtgttt aaggccgtgt cattcaagag ccaggtggtc gcggggacaa actacttcat 180
caaggtgcac gtcggcgacg aggacttcgt acacctgcga gtgttccaat ctctccctca 240
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ccttggggtg gaaggggcan gatctgcact gcttttgcat ttctcttcct aaatttcatt 480
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<210> 76
<211> 666
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<222> 411, 470, 476, 491, 506, 527, 560, 570, 632, 636, 643, 650,
654, 658
<223> n = A, T, C or G
<400> 76
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ttcctggcta ctccatgttg gctagcctct ggtaacctct tacttattat cttcaggaca 240
ctcactacag ggaccaggga tgatgcaaca tccttgtctt tttatgacag gatgtttgct 300
cagettetee aacaataaaa ageaegtggt aaaacaettg eggatattet ggaetgtttt 360
taaaaaatat acagtttacc gaaaatcata ttatcttaca atgaaaagga ntttatagat 420
cagccagtga acaacctttt cccaccatac aaaaattcct tttcccgaan gaaaanggct 480
ttctcaataa ncctcacttt cttaanatct tacaaqatag ccccqanatc ttatcqaaac 540
tcattttagg caaatatgan ttttattgtn cgttacttgt ttcaaaattt ggtattgtga 600
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atatcaatta ccaccccat ctcccatqaa anaaanggqa aanggtqaan ttcntaancg 660
cttaaa
<210> 77
<211> 396
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<213> Homo sapiens
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<221> misc feature
\langle 222 \rangle 31, \overline{5}4, 125, 128, 136, 163, 168, 198
<223> n = A, T, C or G
<400> 77
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catanganta tgccanaata aattccattt ttttgaaaat canctccntg gggctggttt 180
tggtccacag cataacangc actgcctcct tacctgtgag gaatgcaaaa taaagcatgg 240
attaagtgag aagggagact ctcagccttc agcttcctaa attctgtgtc tgtgactttc 300
gaagtttttt aaacctctga atttgtacac atttaaaatt tcaagtgtac tttaaaataa 360
aatacttcta atgggaacaa aaaaaaaaa aaaaaa
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<210> 78
<211> 793
<212> DNA
<213> Homo sapiens
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<221> misc feature
<222> 309, 492, 563, 657, 660, 703, 708, 710, 711, 732, 740, 748,
758, 762, 765, 787
<223> n = A, T, C or G
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taccacagte aaacetggag ecaaaaagga cacaaaggae tetegaceca aactgeecca 180
gaccetetee agaggttggg gtgaccaact catetggact caqacatatg aaqaagetet 240
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tctgtcggct tgaaaattga aaccagaaaa atgtgaaaaa tggctattgt ggaacanatn 660
gacacctgat taggttttgg ttatgttcac cactattttt aanaaaanan nttttaaaat 720
ttggttcaat tntcttttn aaacaatntg tttctacntt gnganctgat ttctaaaaaa 780
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<210> 79
<211> 456
<212> DNA
<213> Homo sapiens
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<222> 89, 195, 255, 263, 266, 286, 353, 384, 423, 425, 436, 441
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gcagctgttg agegeaceta accaetggte atgeececae ecetgetete egcaeceget 180
tecteeegae eecangaeea ggetaettet eeesteetet tgeeteeete etgeeeetge 240
tgcctctgat cgtangaatt gangantgtc ccgccttgtg gctganaatg gacagtggca 300
ggggctggaa atgggtgtgt gtgtgtgtgt gtgtgtgtgt gtgtgtgtgt gcncccccc 360
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<210> 80
<211> 284
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 283
<223> n = A, T, C or G
<400> 80
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gaatagcatg acctccgtgc aaacaggaca agcaaatttg tgatgtgttg attaaaaaga 180
aataaataaa tgtgtatatg tgtaacttgt atgtttatgt ggaatacaga ttgggaaata 240
<210> 81
<211> 671
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 388, 505, 600, 603, 615, 642, 644, 660
<223> n = A, T, C or G
<400> 81
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gaaaggctgg ggatatttgg gttggcttgg ttttgatttt ttgcttgttt gtttgttttg 180
tactaaaaca gtattatctt ttgaatatcg tagggacata agtatataca tgttatccaa 240
tcaagatggc tagaatggtg cetttetgag tgtetaaaac ttgacacccc tggtaaatet 300
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tttcaatgcc gtcattttca gttagatnat tttgcacttt gagattaaaa tgccatgtct 420
atttgattag tettattttt ttattttae aggettatea gteteaetgt tggetgteat 480
tgtgacaaag tcaaataaac ccccnaggac aacacacagt atgggatcac atattgtttg 540
acattaaqct ttqqccaaaa aatqttqcat qtqttttacc tcqacttqct aaatcaatan 600
canaaaggct ggctnataat gttggtggtg aaataattaa tnantaacca aaaaaaaan 660
```

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ormal areas, states, never server, ser
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671
aaaaaaaaaa a
<210> 82
<211> 217
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 35
<223> n = A, T, C or G
<400> 82
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tcttattagg gagttgtatg tcagtgtata aaacatactg tgtggtataa caggcttaat 180
                                                                   217
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<210> 83
<211> 460
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 104, 118, 172, 401, 422, 423, 444, 449
<223> n = A, T, C or G
<400> 83
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aacggagacg caggagaaga acacctgcc gaccaaagag accattgagc angagaagcg 180
gagtgaaatt teetaagate etggaggatt teetaeeeee gteetetteg agaeeeeagt 240
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ctgggcactc cgcgccgatg ccaccggcct gtgggtctct gaagggaccc ccccaatcg 360
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annataaaac acacctcgtg gcancaaana aaaaaaaaaa
<210> 84
<211> 323
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 70, 138, 178, 197, 228, 242, 244, 287, 311
<223> n = A, T, C or G
<400> 84
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gtggtccaan gcattttgct ggcttaacgg gtcccggaac aaaggacacc agctctctaa 120
aattgaagtt tacceganat aacaatettt tgggeagaga tgeetatttt aacaaaenee 180
gtccctgcgc aacaacnaac aatctctggg aaataccggc catgaacntg ctgtctcaat 240
chancatote tetagetgae egateatate gteceagatt actaeanate ataataattg 300
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323
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<210> 85
<211> 771
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 63, 426, 471, 497, 521, 554, 583, 586, 606, 609, 615, 652,
686, 691, 694, 695, 706, 713, 730, 732, 743, 751
<223> n = A, T, C or G
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gaagcaagca actttgactg ctgtcttgga tacacagacc gtattcttca tcctaaattt 180
attgtgggct tcacacggca gctggccaat gaaggctgtg acatcaatgc tatcatcttt 240
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gtgcgtctcc tcagtaaaaa agtcaagaac atgtaaaaac tgtggctttt ctggaatgga 360
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gttatttata gctntaggtt ttctgtgttt aactttttat acnaantttc ctaaactatt 600
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gccacaagct ttttttaaaa aaccantaca nccnngttaa atggtnggtc ccnaatggtt 720
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<210> 86
<211> 628
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 162, 249, 266, 348, 407, 427, 488, 518, 545, 566, 569, 597,
598, 611, 617, 621, 624
<223> n = A, T, C or G
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agttcataca ttcaaagcat ctgaactgta gtttctatag caagccaatt acatccataa 240
gtggagaang aaatagatta atgtcnaagt atgattggtg gagggagcaa ggttgaagat 300
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gaaatattaa tgtttacctt tcaatgtgtg gtatcagctg gactcantaa cacccctttc 420
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teetttenea gtttetgget eetaeeetae tgatttanee agaataagaa aacattttat 540
catchtctgc tttattccca ttaatnaant tttgatgaat aaatctgctt ttatgcnnac 600
                                                                   628
ccaaggaatt nagtggnttc ntcnttgt
<210> 87
<211> 518
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<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> 384, 421, 486
<223> n = A, T, C or G
<400> 87
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agtagtacag ttttaaaatt ttatgcttaa aacaagtttt gtgtaaaaaa tgcagataca 180
ttttacatgg caaatcaatt tttaagtcat cctaaaaatt gattttttt tgaaatttaa 240
aaacacattt aatttcaatt tototottat ataacettta ttactatage atggtttoca 300
ctacagttta acaatgcagc aaaattccca tttcacggta aattgggttt taagcggcaa 360
ggttaaaatg ctttgaggat cctnaatacc ctttgaactt caaatgaagg ttatggttgt 420
naatttaacc ctcatqccat aaqcaqaaqc acaagtttag ctqcattttg ctctaaactg 480
                                                                 518
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<210> 88
<211> 1844
<212> DNA
<213> Homo sapiens
<400> 88
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ttccatcttc ttggtgctgg gaagccatat atgtgtcttt tactcaagct aaggggtata 240
agettatgtg ttgaatttgc tacatetata tttcacatat tetcacaata agagaatttt 300
gaaatagaaa tatcatagaa catttaagaa agtttagtat aaataatatt ttgtgtgttt 360
taatcccttt gaagggatct atccaaagaa aatattttac actgagctcc ttcctacacg 420
totcagtaac agatoctgtg ttagtotttg aaaatagoto attttttaaa tgtcagtgag 480
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acatagggca atctgtgaat atgtattata agcagcattc cagaaaagta gttggtgaaa 660
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atttgaagtt caaaggtgta ttcaggatcc tcaaagcatt ttaaccttgc cgcttaaaac 1500
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aaaatcaatc tttaggatga cttaaaaatt gatttgccat gtaaaatgta tctgcatttt 1680
ttacacaaaa cttqttttaa qcataaaatt ttaaaactqt actacttqat qtattataca 1740
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THEOR ATTEN AREA THEOR ATTENDED TO ATTEND AT A TO ATT ANY ATTENDED TO A TO ATTEND ATTENDED TO ATTENDED TO ATTENDED TO ATTENDED TO ATTENDED ATTENDED
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ttttqaacca tatqtattaa accataaaca gtataatgtt gttataataa aacaggcaat 1800
aaatttataa ataaaagctq aaaaaaaaaaa aaaaaaaaaa aaaa
<210> 89
<211> 523
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 288, 352, 369, 398, 475, 511, 513
<223> n = A, T, C or G
<400> 89
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acaatatgat gtagaaaatg ctaagccaga gatatagaaa ggtcctattg ggtccttctg 180
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gccctggcat gacttgaacc caaccacaga ctgggaaagg gagcctttcg anagtggatc 360
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ggtgctcaag aaaagtttgc agaatggata aatgaaggat caagggaatt aatanatgaa 480
                                                                   523
taattgaatg gtggctcaat aagaatgact ncnttgaatg acc
<210> 90
<211> 604
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 563
<223> n = A, T, C or G
<400> 90
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qcaaaqqaaa tagccaatat gtqtcqtttc tatqaaatga agccagaccg agatgtcaat 120
ctcacccacc aactaaatcc caaagtcaaa agcttcagcc agtttatctc agagaaccag 180
gggagcette aagggeatgt agaaaateag etgtteagat aggeetetge accacacage 240
ctctttcctc tctgatcctt ttcctcttta cggcacaaca ttcatgtttg acagaacatg 300
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cactgcattg gtgataggac acggtaattt gattcacatt taacttgcta gttagtgata 420
aggggtggta cacctgtttg gtaaaatgag aagcctcgga aacttgggag cttctctcct 480
accactaatg gggagggcag attattactg ggatttctcc tggggtgaat taatttcaag 540
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                                                                   604
CCCC
<210> 91
<211> 858
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
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<222> 570, 591, 655, 664, 667, 683, 711, 759, 760, 765, 777, 787,
792, 794, 801, 804, 809, 817, 820
<223> n = A, T, C or G
<400> 91
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cattteetta tteaageeat gettttetgt gatattetga teetagttga acatacagaa 180
ataaatgtct aaaacagcac ctcgattctc gtctataaca ggactaagtt cactgtgatc 240
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geoeggtace caattegeee tatagtgagt egtattaege gegeteactg geogtegttt 480
tacaacqtcq tqactqqqaa aaccctqqcq ttacccaact taatcqcctt qcaqcacatc 540
cccctttcgc cagctggcgt aatagcgaan agcccgcacc gatcgccctt ncaacagttg 600
cqcaqcctqa atqqcqaatq qqacqcqccc tqtaqcqqcq cattaaaqcq cqgcnqqqtq 660
tggnggntce cecaegtgae cgntacaett ggeagegeet taegeeggte nttegettte 720
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cctttanggg tncnaattaa nggnttacng gaccttngan cccaaaaact ttgattaggg 840
ggaaggtccc cgaagggg
<210> 92
<211> 585
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 317, 319, 320, 321, 325, 327, 328, 330, 331, 332, 460, 462,
483, 485, 487, 523, 538, 566, 584
<223> n = A, T, C or G
<400> 92
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atatcacaga aaagcatggc ttgaataagg aaatgacaat tttttccact tatctgatca 240
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aaaaaataat aatcatnann naaanannan nngaagggcg gccgccaccg cggtggagct 360
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gentnangtg taaaageetg ggggtgeeta attgagtgag etnaeteaca ttaattgngt 540
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<210> 93
<211> 567
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 82, 158, 230, 232, 253, 266, 267, 268, 269, 270, 271, 272,
273, 274, 275, 276, 277, 278, 279, 280, 281, 282, 283, 284,
285, 286, 287, 295, 303, 307, 314, 349, 352, 354, 356, 366,
```

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369, 379, 382, 386, 393, 404, 427, 428, 446, 450, 452
<223> n = A, T, C or G
<221> misc feature
<222> 453, 454, 459, 462, 480, 481, 483, 488, 493, 501, 509, 511,
512, 518, 520, 525, 526, 532, 541, 557
<223> n = A, T, C or G
<400> 93
cggcagtgtt gctgtctgcg tgtccacctt ggaatctggc tgaactggct gggaggacca 60
agactgcggc tggggtgggc anggaaggga accgggggct gctgtgaagg atcttggaac 120
ttccctgtac ccaccttccc cttgcttcat gtttgtanag gaaccttgtg ccggccaagc 180
ccagtttcct tgtgtgatac actaatgtat ttgctttttt tgggaaatan anaaaaatca 240
attaaattgc tantgtttct ttgaannnnn nnnnnnnnn nnnnnnngg ggggncgccc 300
concedenge aacheecect titgticect tiaatigaaa gettaating enenentege 360
gttaancent gggecaaane tngttneeeg tgntgaaatt gttnateeee teecaaatte 420
cccccnncc ttccaaaccc ggaaancctn annntgttna ancccggggg gttgcctaan 480
ngnaattnaa cenaaceee ntttaaatng nntttgenen ceaenngeee enettteeea 540
nttcggggaa aaccctntcc gtgccca
                                                                   567
<210> 94
<211> 620
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 169, 171, 222, 472, 528, 559, 599
<223> n = A, T, C or G
<400> 94
actagtcaaa aatgctaaaa taatttggga gaaaatattt tttaagtagt gttatagttt 60
catgtttatc ttttattatg ttttgtgaag ttgtgtcttt tcactaatta cctatactat 120
gccaatattt ccttatatct atccataaca tttatactac atttgtaana naatatgcac 180
gtgaaactta acactttata aggtaaaaat gaggtttcca anatttaata atctgatcaa 240
gttcttgtta tttccaaata qaatggactt ggtctgttaa gggctaagga gaagaggaag 300
ataaggttaa aagttgttaa tgaccaaaca ttctaaaaga aatgcaaaaa aaaagtttat 360
tttcaageet tegaaetatt taaggaaage aaaateattt eetaaatgea tateatttgt 420
gagaatttet cattaatate etgaateatt cattteacta aggeteatgt tnacteegat 480
atgtetetaa gaaagtaeta ttteatggte caaacetggt tgecatantt gggtaaagge 540
tttcccttaa gtgtgaaant atttaaaatg aaattttcct ctttttaaaa attctttana 600
                                                                   620
agggttaagg gtgttgggga
<210> 95
<211> 470
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 61, 67, 79, 89, 106, 213, 271, 281, 330, 354, 387, 432, 448
<223> n = A, T, C or G
<400> 95
```

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cteqacette tetgeacage ggatgaacee tgagcagetg aagaceagaa aageeactat 60
nactttntgc ttaattcang agcttacang attcttcaaa gagtgngtcc agcatccttt 120
gaaacatgag ttcttaccag cagaagcaga cctttacccc accacctcag cttcaacagc 180
agcaggtgaa acaacccatc cagcctccac ctnaggaaat atttgttccc acaaccaagg 240
agccatgcca ctcaaaggtt ccacaacctg naaacacaaa nattccagag ccaggctgta 300
ccaaggtccc tgagccaggg ctgtaccaan gtccctgagc caggttgtac caangtccct 360
gagccaggat gtaccaaggt ccctgancca ggttgtccaa ggtccctgag ccaggctaca 420
ccaagggect gngccaggea gcatcaangt ccctgaccaa ggcttatcaa
<210> 96
<211> 660
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 299, 311, 360, 426, 538, 540, 542, 553, 563, 565, 592, 603,
604, 618, 633, 647, 649, 651, 653
<223> n = A, T, C or G
<400> 96
ttttttttt tttttttt ggaattaaaa gcaatttaat gagggcagag caggaaacat 60
gcatttettt teattegaat etteagatga accetgagea geegaagaee agaaaageea 120
tgaagacttt ctgcttaatt caggggctta caggattctt cagagtgtgt gtgaacaaaa 180
gctttatagt acgtattttt aggatacaaa taagagagag actatggctt ggggtgagaa 240
tgtactgatt acaaggteta cagacaatta agacacagaa acagatggga agagggtgnc 300
cagcatctgg nggttggett ctcaaggget tgtctgtgca ccaaattact tctgcttggn 360
cttctqctqa qctqqqcctq qaqtqaccqt tqaaqqacat qqctctqqta cctttqtqta 420
gcctgncaca ggaactttgg tgtatccttg ctcaggaact ttgatggcac ctggctcagg 480
aaacttgatg aagcettggt caagggacct tgatgettge tggetcaggg accttggngn 540
ancetggget canggacett tgneneaace ttggetteaa gggaceettg gnacateetg 600
gennagggae cettgggnee aaccetggge ttnagggaee etttggntne nancettgge 660
<210> 97
<211> 441
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 12, 308
<223> n = A, T, C or G
<400> 97
gggaccatac anagtattcc tctcttcaca ccaggaccag ccactgttgc agcatgagtt 60
cccagcagca gaagcagccc tgcatcccac cccctcagct tcagcagcag caggtgaaac 120
agocttgoca geetecaeet caggaaceat geateceeaa aaceaaggag eeetgeeaee 180
ccaaggtgcc tgagccctgc caccccaaag tgcctgagcc ctgccagccc aaggttccag 240
agecatgeca ecceaaggtg ectgageet geetteaat agteacteea geaceageee 300
agcagaanac caagcagaag taatgtggtc cacagccatg cccttgagga gccggccacc 360
agatgctgaa tcccctatcc cattctgtgt atgagtccca tttgccttgc aattagcatt 420
ctgtctcccc caaaaaaaaa a
                                                                   441
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there are some other area, are and a property of the property
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<210> 98
<211> 600
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> 295, 349, 489, 496, 583
<223> n = A, T, C or G
<400> 98
gtatteetet etteacacca ggaccageca etgttgeage atgagtteee ageageagaa 60
gcagccetge ateccaecee etcagettea gcagcageag gtgaaacage ettgecagee 120
tocaceteag gaaceatgea teeceaaaae caaggageee tgeeaeeeea aggtgeetga 180
gccctgccac cccaaaqtqc ctqaqccctq ccaqcccaaq qttccaqaqc catqccaccc 240
caaggtgeet gageeetgee etteaatagt caeteeagea eeageeeage agaanaceaa 300
gcagaagtaa tgtggtccac agccatgccc ttgaggagcc ggccaccana tgctgaatcc 360
cetateceat tetgtgtatg agteceattt geettgeaat tageattetg teteceecaa 420
aaaagaatgt getatgaage tttettteet acacactetg agtetetgaa tgaagetgaa 480
ggtcttaant acaganctag ttttcagctg ctcagaattc tctgaagaaa agatttaaga 540
tgaaaggcaa atgattcagc tccttattac cccattaaat tcnctttcaa ttccaaaaaa 600
<210> 99
<211> 667
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 345, 562, 635
<223> n = A, T, C or G
<400> 99
actagtgact gagttcctgg caaagaaatt tgacctggac cagttgataa ctcatgtttt 60
accatttaaa aaaatcagtg aaggatttga gctgctcaat tcaggacaaa gcattcgaac 120
ggtcctgacg ttttgagatc caaagtggca ggaggtctgt gttgtcatgg tgaactggag 180
tttctcttgt gagagttccc tcatctgaaa tcatgtatct gtctcacaaa tacaagcata 240
agtagaagat ttgttgaaga catagaaccc ttataaagaa ttattaacct ttataaacat 300
ttaaagtctt gtgagcacct gggaattagt ataataacaa tgttnatatt tttgatttac 360
attttgtaag getataattg tatettttaa gaaaacatac ettggattte tatgttgaaa 420
tggagatttt taagagtttt aaccagctgc tgcagatata ttactcaaaa cagatatagc 480
gtataaagat atagtaaatg catctcctag agtaatattc acttaacaca ttggaaacta 540
ttatttttta gatttgaata tnaatgttat tttttaaaca cttgttatga gttacttggg 600
attacatttt gaaatcagtt cattccatga tgcanattac tgggattaga ttaagaaaga 660
cggaaaa
                                                                   667
<210> 100
<211> 583
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
```

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<222> 404, 506, 514, 527, 528, 538, 548, 556, 568, 569
<223> n = A, T, C or G
<400> 100
gttttgtttg taagatgatc acagtcatgt tacactgatc taaaggacat atatataacc 60
ctttaaaaaa aaaatcactg cctcattctt atttcaagat gaatttctat acagactaga 120
tgtttttctg aagatcaatt agacattttg aaaatgattt aaagtgtttt ccttaatgtt 180
ctctgaaaac aagtttcttt tgtagtttta accaaaaaag tgcccttttt gtcactggat 240
tctcctagca ttcatgattt ttttttcata caatgaaatt aaaattgcta aaatcatgga 300
ctggctttct ggttggattt caggtaagat gtgtttaagg ccagagcttt tctcagtatt 360
tgattttttt ccccaatatt tgatttttta aaaatataca catnggtgct gcatttatat 420
ctgctqqttt aaaattctqt catatttcac ttctaqcctt ttaqttatqq caaatcatat 480
tttactttta cttaaagcat ttggtnattt ggantatctg gttctannct aaaaaaanta 540
                                                                   583
attctatnaa ttgaantttt ggtactcnnc catatttgga tcc
<210> 101
<211> 592
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 218, 497, 502, 533, 544, 546, 548, 550, 555
<223> n = A, T, C or G
<400> 101
gtggagacgt acaaagagca gccgctcaag acacctggga agaaaaagaa aggcaagccc 60
gggaaacgca aggagcagga aaagaaaaaa cggcgaactc gctctgcctg gttagactct 120
ggagtgactg ggagtgggct agaaggggac cacctgtctg acacctccac aacgtcgctg 180
gagetegatt caeggaggea ttgaaatttt cageaganae ettecaagga catattgeag 240
qattctqtaa taqtqaacat atqqaaaqta ttaqaaatat ttattqtctq taaatactqt 300
aaatgcattq qaataaaact qtctccccca ttqctctatq aaactqcaca ttqqtcattq 360
tgaatatttt tttttttgcc aaggctaatc caattattat tatcacattt accataattt 420
attttgtcca ttgatgtatt tattttgtaa atgtatcttg gtgctgctga atttctatat 480
tttttgtaca taatgcnttt anatatacct atcaagtttg ttgataaatg acncaatgaa 540
gtgncncnan ttggnggttg aatttaatga atgcctaatt ttattatccc aa
<210> 102
<211> 587
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 91, 131, 256, 263, 332, 392, 400, 403, 461, 496, 497, 499,
510, 511, 518, 519, 539, 554, 560, 576
<223> n = A, T, C or G
<400> 102
egtectaage acttagacta cateagggaa gaacacagae cacatecetg teetcatgeg 60
gettatgttt tetggaagaa agtggagaee nagteettgg etttaggget eeceggetgg 120
gggctgtgca ntccggtcag ggcgggaagg gaaatgcacc gctgcatgtg aacttacagc 180
ccaggeggat geocettece ttageactae etggeetect geatececte geoteatgtt 240
cctcccacct tcaaanaatg aanaacccca tgggcccagc cccttgccct ggggaaccaa 300
```

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ggcagccttc caaaactcag gggctgaagc anactattag ggcaggggct gactttgggt 360
gacactgece attecetete agggeagete angteaecen ggnetettga acceageetg 420
ttcctttgaa aaaqgqcaaa actgaaaagg gcttttccta naaaaagaaa aaccagggaa 480
ctttgccagg gcttcnntnt taccaaaacn ncttctcnng gatttttaat tccccattng 540
gcctccactt accnggggcn atgccccaaa attaanaatt tcccatc
<210> 103
<211> 496
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
\langle 222 \rangle 2, 1\overline{7}, 66, 74, 82, 119, 164, 166, 172, 200, 203, 228, 232,
271, 273, 415, 423, 445, 446, 473
<223> n = A, T, C or G
<400> 103
anaggactgg ccctacntgc tetetetegt cetacetate aatgeceaae atggcagaac 60
ctgcanccct tggncactgc anatggaaac ctctcagtgt cttgacatca ccctacccnt 120
qcqqtqqqtc tccaccacaa ccactttqac tctqtqqtcc ctqnangqtq qnttctcctq 180
actggcagga tggaccttan ccnacatatc cctctgttcc ctctgctnag anaaaqaatt 240
cccttaacat gatataatcc acccatgcaa ntngctactg gcccagctac catttaccat 300
ttgcctacag aatttcattc agtctacact ttggcattct ctctggcgat agagtgtggc 360
tgggctgacc gcaaaaggtg ccttacacac tggcccccac cctcaaccgt tgacncatca 420
gangettgee teeteettet gattnnceee eatgttggat ateagggtge tenagggatt 480
                                                                 496
ggaaaagaaa caaaac
<210> 104
<211> 575
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 18, 19, 45, 68, 77, 132, 155, 174, 219, 226, 238, 259, 263,
271, 273, 306, 323, 339, 363, 368, 370, 378, 381, 382, 436,
440, 449, 450, 456, 481, 485, 496, 503, 510, 512, 515, 528,
542, 552
<223> n = A, T, C or G
<400> 104
ctatggangt ggtttenggg gtggetettg ceaactggga agaageegtg gtgtetetae 120
ctgttcaact cngtttgtgt ctgggggatc aactnggggc tatggaagcg gctnaactgt 180
tgttttggtg gaagggctgg taattggctt tgggaagtng cttatngaag ttggcctngg 240
gaagttgcta ttgaaagtng ccntggaagt ngntttggtg gggggttttg ctggtggcct 300
ttgttnaatt tgggtgcttt gtnaatggcg gcccctcnc ctgggcaatg aaaaaaatca 360
conatgongn aaacctonac nnaacagoot gggottooot cacotogaaa aaagttgoto 420
ccccccaaa aaaggncaan cccctcaann tggaangttg aaaaaatcct cgaatgggga 480
ncccnaaaac aaaaancccc contttcocn gnaanggggg aaataccncc cccccactta 540
                                                                 575
cnaaaaccct tntaaaaaac cccccgggaa aaaaa
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<211> 619
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> 260, 527, 560, 564, 566, 585, 599
<223> n = A, T, C or G
<400> 105
cactagtagg atagaaacac tgtgtcccga gagtaaggag agaagctact attgattaga 60
qcctaaccca qqttaactqc aaqaaqaqqc qqqatacttt caqctttcca tqtaactqta 120
tgcataaagc caatgtagtc cagtttctaa gatcatgttc caagctaact gaatcccact 180
tcaatacaca ctcatgaact cctgatggaa caataacagg cccaagcctg tggtatgatg 240
tgcacacttg ctagactcan aaaaaatact actctcataa atgggtggga gtattttggt 300
gacaacctac tttgcttggc tgagtgaagg aatgatattc atatattcat ttattccatg 360
qacatttaqt taqtqctttt tatataccaq qcatqatqct qaqtqacact cttqtqtata 420
tttccaaatt tttqtacaqt cqctqcacat atttqaaatc atatattaag acttccaaaa 480
aatgaagtee etggttttte atggeaactt gateagtaaa ggatteneet etgtttggta 540
cttaaaacat ctactatatn gttnanatga aatteetttt cecencetee egaaaaaana 600
                                                                    619
aagtggtggg gaaaaaaaa
<210> 106
<211> 506
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
\langle 222 \rangle 8, 2\overline{1}, 31, 32, 58, 75, 89, 96, 99, 103, 122, 126, 147, 150,
158, 195, 210, 212, 219, 226, 246, 248, 249, 255, 258, 261,
263, 265, 275, 304, 317, 321, 331, 337, 340, 358, 371, 377,
380, 396, 450, 491
<223> n = A, T, C or G
<400> 106
cattggtnct ttcatttgct ntggaagtgt nnatctctaa cagtggacaa agttcccngt 60
gccttaaact ctgtnacact tttgggaant gaaaanttng tantatgata ggttattctg 120
angtanagat gttctggata ccattanatn tgcccccngt gtcagaggct catattgtgt 180
tatgtaaatg gtatntcatt cgctactatn antcaattng aaatanggtc tttgggttat 240
gaatantnng cageneanet nanangetgt etgtngtatt cattgtggte atageacete 300
acancattgt aacctcnatc nagtgagaca nactagnaan ttcctagtga tggctcanga 360
ttccaaatgg nctcatntcn aatgtttaaa agttanttaa gtgtaagaaa tacagactgg 420
atgttccacc aactagtacc tgtaatgacn ggcctgtccc aacacatctc ccttttccat 480
                                                                    506
gactgtggta necegcateg gaaaaa
<210> 107
<211> 452
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 289, 317, 378
```

112

113

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<223> n = A, T, C or G
<400> 107
gttgagtctg tactaaacag taagatatct caatgaacca taaattcaac tttgtaaaaa 60
tcttttgaag catagataat attgtttggt aaatgtttct tttgtttggt aaatgtttct 120
tttaaagacc ctcctattct ataaaactct gcatgtagag gcttgtttac ctttctctct 180
ctaaggttta caataggagt ggtgatttga aaaatataaa attatgagat tggttttcct 240
gtggcataaa ttgcatcact gtatcatttt cttttttaac cggtaagant ttcagtttgt 300
tggaaagtaa ctgtganaac ccagtttccc gtccatctcc cttagggact acccatagaa 360
catgaaaagg tccccacnga agcaagaaga taagtctttc atggctgctg gttgcttaaa 420
                                                                   452
ccactttaaa accaaaaaat tccccttgga aa
<210> 108
<211> 502
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 22, 31, 126, 168, 183, 205, 219, 231, 236, 259, 283, 295,
296, 298, 301, 340, 354, 378, 383, 409, 433, 446, 455, 466,
<223> n = A, T, C or G
<400> 108
atcttcttcc cttaattagt tnttatttat ntattaaatt ttattgcatg tcctggcaaa 60
caaaaagaga ttgtagattg gcttctggct ccccaaaagc ccataacaga aagtaccaca 120
agaccncaac tgaagcttaa aaaatctatc acatgtataa tacctttnga agaacattaa 180
tanagcatat aaaactttta acatntgctt aatgttgtnc aattataaaa ntaatngaaa 240
aaaatqtccc tttaacatnc aatatcccac ataqtqttat ttnaqqqqat taccnnqnaa 300
naaaaaaaqq qtaqaaqqqa tttaatqaaa actctqcttn ccatttctqt ttanaaacqt 360
ctccagaaca aaaacttntc aantctttca gctaaccgca tttgagctna ggccactcaa 420
aaactccatt agncccactt tctaanggte tctanagctt actaancctt ttgaccctt 480
                                                                   502
accetggnta etcetgeeet ca
<210> 109
<211> 1308
<212> DNA
<213> Homo sapiens
<400> 109
accogaggte tegetaaaat cateatggat teacttggeg eegteageae tegacttggg 60
tttgatettt teaaagaget gaagaaaaca aatgatggea acatettett tteeeettgtg 120
ggcatcttqa ctgcaattgg catggtcctc ctggggaccc gaggagccac cgcttcccag 180
ttggaggagg tgtttcactc tgaaaaagag acgaagagct caagaataaa ggctgaagaa 240
aaagaggtga ttgagaacac agaagcagta catcaacaat tccaaaagtt tttgactgaa 300
ataagcaaac tcactaatga ttatgaactg aacataacca acaggctgtt tggagaaaaa 360
acatacetet teetteaaaa ataettagat tatgttgaaa aatattatea tgeatetetg 420
gaacctgttg attttgtaaa tgcagccgat gaaagtcgaa agaagattaa ttcctgggtt 480
gaaagcaaaa caaatgaaaa aatcaaggac ttgttcccag atggctctat tagtagctct 540
accaagetgg tgetggtgaa catggtttat tttaaaggge aatgggacag ggagtttaag 600
aaagaaaata ctaaggaaga gaaattttgg atgaataaga gcacaagtaa atctgtacag 660
atgatgacac agagecatte etttagette aettteetgg aggaettgea ggecaaaatt 720
ctagggattc catataaaaa caacgaccta agcatgtttg tgcttctgcc caacgacatc 780
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gatggcctgg agaagataat agataaaata agtcctgaga aattggtaga gtggactagt 840
ccagggcata tggaagaaag aaaggtgaat ctgcacttgc cccggtttga ggtggaggac 900
agttacgatc tagaggcggt cctggctgcc atggggatgg gcgatgcctt cagtgagcac 960
aaagccgact actcgggaat gtcgtcaggc tccgggttgt acgcccagaa gttcctgcac 1020
agttcctttg tggcagtaac tgaggaaggc accgaggctg cagctgccac tggcataggc 1080
tttactgtca catcogcccc aggtcatgaa aatgttcact gcaatcatcc cttcctgttc 1140
ttcatcaggc acaatgaatc caacagcatc ctcttcttcg gcagattttc ttctccttaa 1200
gatgategtt gecatggeat tgetgetttt ageaaaaaac aactaceagt gttacteata 1260
tgattatgaa aatcgtccat tcttttaaat ggtggctcac ttgcattt
<210> 110
<211> 391
<212> PRT
<213> Homo sapiens
<400> 110
Met Asp Ser Leu Gly Ala Val Ser Thr Arg Leu Gly Phe Asp Leu Phe
                                    10
Lys Glu Leu Lys Lys Thr Asn Asp Gly Asn Ile Phe Phe Ser Pro Val
            20
                                25
Gly Ile Leu Thr Ala Ile Gly Met Val Leu Leu Gly Thr Arg Gly Ala
                            40
Thr Ala Ser Gln Leu Glu Glu Val Phe His Ser Glu Lys Glu Thr Lys
                        55
                                             60
Ser Ser Arg Ile Lys Ala Glu Glu Lys Glu Val Ile Glu Asn Thr Glu
                    70
                                         75
Ala Val His Gln Gln Phe Gln Lys Phe Leu Thr Glu Ile Ser Lys Leu
                85
                                    90
Thr Asn Asp Tyr Glu Leu Asn Ile Thr Asn Arg Leu Phe Gly Glu Lys
            100
                                105
                                                     110
Thr Tyr Leu Phe Leu Gln Lys Tyr Leu Asp Tyr Val Glu Lys Tyr Tyr
        115
                            120
                                                 125
His Ala Ser Leu Glu Pro Val Asp Phe Val Asn Ala Ala Asp Glu Ser
                        135
                                             140
Arg Lys Lys Ile Asn Ser Trp Val Glu Ser Lys Thr Asn Glu Lys Ile
                    150
                                        155
Lys Asp Leu Phe Pro Asp Gly Ser Ile Ser Ser Ser Thr Lys Leu Val
                                    170
Leu Val Asn Met Val Tyr Phe Lys Gly Gln Trp Asp Arg Glu Phe Lys
                                185
                                                     190
Lys Glu Asn Thr Lys Glu Glu Lys Phe Trp Met Asn Lys Ser Thr Ser
                            200
                                                 205
Lys Ser Val Gln Met Met Thr Gln Ser His Ser Phe Ser Phe Thr Phe
    210
                        215
                                             220
Leu Glu Asp Leu Gln Ala Lys Ile Leu Gly Ile Pro Tyr Lys Asn Asn
                    230
                                         235
Asp Leu Ser Met Phe Val Leu Leu Pro Asn Asp Ile Asp Gly Leu Glu
                245
                                     250
                                                         255
Lys Ile Ile Asp Lys Ile Ser Pro Glu Lys Leu Val Glu Trp Thr Ser
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TIIT	50	Ser	GIII	TIGU	GIU	55	val	1116	1112	Set	60	цуз	Giu	1111	цуз
Sar		Δra	Tle	Lus	Δla		Glu	T.vs	Glu	Val		Ara	Tle	T.VS	Δla
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Cid	OTY	шуо	OIG	85	014	71011		014	90	vai	1110	0111	01	95	0111
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Lys	Gly		Trp	Asp	Arg	Glu	Phe	Lys	Lys	Glu	Asn		Lys	GLu	Glu
-	D1	195			-	0	200	0 -	Ŧ	0	11. 7	205	M - +	N/ - L	m)
Lys		Trp	Met	Asn	Lys		Thr	ser	ьуs	ser	220	GIN	мет	мет	TUL
Cln	210	шіс	Sor	Dho	Car	215 Pho	Thr	Dho	LOU	Clu		Ton	Gln	ΛΙο	Twe
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His	Lys	Ala	Asp	-	Ser	Gly	Met	Ser		Gly	Ser	Gly	Leu	_	Ala
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Gln	Lys	Phe		His	Ser	Ser	Phe		Ala	Val	Thr	GLu		GLY	Thr
C3	7.7.	70.1 -	340	71 -	mla .a	C1	Tla	345	Dh.a	mb so	77.0.7	mb~	350	70.7	D 20.0
GIU	Ala	355		Ата	1111.	СТУ	Ile 360	СТУ	Pne	1111	Va⊥	365	ser	Ald	PIO
Clar	шіс			V-1	uic	Circ	Asn	Hic	Dro	Dha	Tou		Dha	Tla	7\ r \
σтλ	370	GIU	W211	val	11172	375		1112	110	r 11G	380	T 11G	1116	TTG	лгу
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100

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			740					745					Phe 750		
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Lys Ser Ile Gln Asp Leu Arg Arg Phe Phe Leu His His Leu Ile
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Ser Ala Trp Leu Asp Ser Gly Val Thr Gly Ser Gly Leu Glu Gly Asp
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His

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Ser Gln Glu Gly Gly Ser Gly Ser Tyr Glu Glu Gly Cys Gln Ser
                               105
           100
Leu Met Glu Tyr Ala Trp Gly Arg Ala Ala Ala Ala Met Leu Phe Cys
                                               125
                           120
Gly Phe Ile Ile Leu Val Ile Cys Phe Ile Leu Ser Phe Phe Ala Leu
                       135
Cys Gly Pro Gln Met Leu Val Phe Leu Arg Val Ile Gly Gly Leu Leu
                   150
                                       155
Ala Leu Ala Ala Val Phe Gln Ile Ile Ser Leu Val Ile Tyr Pro Val
                                   170
                                                       175
               165
Lys Tyr Thr Gln Thr Phe Thr Leu His Ala Asn Pro Ala Val Thr Tyr
                               185
Ile Tyr Asn Trp Ala Tyr Gly Phe Gly Trp Ala Ala Thr Ile Ile Leu
       195
                           200
                                               205
Ile Gly Cys Ala Phe Phe Cys Cys Leu Pro Asn Tyr Glu Asp Asp
   210
                       215
                                           220
Leu Leu Gly Asn Ala Lys Pro Arg Tyr Phe Tyr Thr Ser Ala
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                                       235
<210> 175
<211> 4181
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 3347, 3502, 3506, 3520, 3538, 3549, 3646, 3940, 3968, 3974,
4036, 4056, 4062, 4080, 4088, 4115
<223> n = A, T, C or G
<400> 175
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actetteaca atgaacaaac tgtatategg aaaceteage gagaacgeeg eeceetegga 300
cctagaaagt atcttcaagg acgccaagat cccggtgtcg ggacccttcc tggtgaagac 360
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qcaaaggatt cggaaacttc agatacgaaa tatcccgcct catttacagt gggaggtgct 540
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ggaaactgca gttgtaaatg taacctattc cagtaaggac caagctagac aagcactaga 660
caaactgaat ggatttcagt tagagaattt caccttgaaa gtagcctata tccctgatga 720
aatggccgcc cagcaaaacc ccttgcagca gccccgaggt cgccgggggc ttgggcagag 780
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tctgcgcctg ctggttccca cccaatttgt tggagccatc ataggaaaag aaggtgccac 900
catteggaac atcaccaaac agacceagte taaaategat gteeacegta aagaaaatge 960
gggggetget gagaagtega ttactateet etetaeteet gaaggeaeet etgeggettg 1020
taaqtctatt ctqqaqatta tqcataaqga aqctcaaqat ataaaattca cagaaqagat 1080
ccccttgaag attttagctc ataataactt tgttggacgt cttattggta aagaaggaag 1140
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aaatettaaa aaaattgage aagacacaga cactaaaate acgatatete cattgcagga 1200 attgacgctg tataatccag aacgcactat tacagttaaa ggcaatgttg agacatgtgc 1260 caaagctgag gaggagatca tgaagaaaat cagggagtct tatgaaaatg atattgcttc 1320 tatgaatctt caagcacatt taattoctgg attaaatctg aacgccttgg gtctgttccc 1380 acceaettea gggatgeeae eteceaeete agggeeeeet teageeatga eteeteeeta 1440 cccgcagttt gagcaatcag aaacggagac tgttcatcag tttatcccag ctctatcagt 1500 cggtgccatc atcggcaagc agggccagca catcaagcag ctttctcgct ttgctggagc 1560 ttcaattaag attgctccag cggaagcacc agatgctaaa gtgaggatgg tgattatcac 1620 tggaccacca gaggeteagt teaaggetea gggaagaatt tatggaaaaa ttaaagaaga 1680 aaactttgtt agtcctaaag aagaggtgaa acttgaagct catatcagag tgccatcctt 1740 tgctgctggc agagttattg gaaaaggagg caaaacggtg aatgaacttc agaatttgtc 1800 aagtgcagaa gttgttgtcc ctcgtgacca gacacctgat gagaatgacc aagtggttgt 1860 caaaataact ggtcacttct atgcttgcca ggttgcccag agaaaaattc aggaaattct 1920 gactcaggta aagcagcacc aacaacagaa ggctctgcaa agtggaccac ctcagtcaag 1980 acqqaaqtaa aqqctcaqqa aacaqcccac cacaqaqqca qatqccaaac caaaqacaqa 2040 ttgcttaacc aacagatggg cgctgacccc ctatccagaa tcacatgcac aagtttttac 2100 ctagccagtt gtttctgagg accaggcaac ttttgaactc ctgtctctgt gagaatgtat 2160 actttatget etetgaaatg tatgacacce agetttaaaa caaacaaaca aacaaacaaa 2220 aaaagggtgg gggagggagg gaaagagaag agctctgcac ttccctttgt tgtagtctca 2280 cagtataaca gatattotaa ttottottaa tattoococa taatgocaga aattggotta 2340 atgatgettt cactaaatte atcaaataga ttgeteetaa atceaattgt taaaattgga 2400 tcagaataat tatcacagga acttaaatgt taagccatta gcatagaaaa actgttctca 2460 qttttatttt tacctaacac taacatqaqt aacctaaqqq aaqtqctqaa tqqtqttqqc 2520 aggggtatta aacgtgcatt tttactcaac tacctcaggt attcagtaat acaatgaaaa 2580 gcaaaattgt tcctttttt tgaaaatttt atatacttta taatgataga agtccaaccg 2640 ttttttaaaa aataaattta aaatttaaca gcaatcagct aacaggcaaa ttaagatttt 2700 tacttctggc tggtgacagt aaagctggaa aattaatttc agggtttttt gaggcttttg 2760 acacagttat tagttaaatc aaatgttcaa aaatacggag cagtgcctag tatctggaga 2820 gcagcactac catttattct ttcatttata gttgggaaag tttttgacgg tactaacaaa 2880 qtqqtcqcaq qaqattttqq aacqqctqqt ttaaatqqct tcaqqaqact tcaqtttttt 2940 gtttagetae atgattgaat geataataaa tgetttgtge ttetgaetat eaataeetaa 3000 agaaagtgca tcagtgaaga gatgcaagac tttcaactga ctggcaaaaa gcaagcttta 3060 gcttgtctta taggatgctt agtttgccac tacacttcag accaatggga cagtcataga 3120 tggtgtgaca gtgtttaaac gcaacaaaag gctacatttc catggggcca gcactgtcat 3180 qaqcctcact aagctatttt qaaqattttt aagcactgat aaattaaaaa aaaaaaaaa 3240 aaattagact ccaccttaag tagtaaagta taacaggatt tctgtatact gtgcaatcag 3300 ttctttqaaa aaaaaqtcaa aaqataqaga atacaagaaa aqttttnggg atataatttq 3360 aatgactgtg aaaacatatg acctttgata acgaactcat ttgctcactc cttgacagca 3420 aagcccagta cqtacaattg tqttqqqtqt qqqtqqtctc caagqccacq ctqctctctg 3480 aattgatttt ttgagttttg gnttgnaaga tgatcacagn catgttacac tgatcttnaa 3540 ggacatatnt tataaccett taaaaaaaaa atcccctgcc tcattcttat ttcgagatga 3600 atttcgatac agactagatg tctttctgaa gatcaattag acattntgaa aatgatttaa 3660 agtgttttcc ttaatgttct ctgaaaacaa gtttcttttg tagttttaac caaaaaagtg 3720 ccctttttgt cactggtttc tcctagcatt catgattttt ttttcacaca atgaattaaa 3780 attgctaaaa tcatggactg gctttctggt tggatttcag gtaagatgtg tttaaggcca 3840 gagettttet eagtatttga tttttteee eaatatttga ttttttaaaa atatacacat 3900 aggagetgea tttaaaacet getggtttaa attetgtean attteaette tageetttta 3960 gtatggcnaa tcanaattta cttttactta agcatttgta atttggagta tctggtacta 4020 gctaagaaat aattenataa ttgagttttg tactenecaa anatgggtca tteeteatgn 4080 ataatgtncc cccaatgcag cttcattttc caganacctt gacgcaggat aaattttttc 4140 4181 atcatttagg tccccaaaaa aaaaaaaaaa a

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Ala Lys Ala Glu Glu Glu Ile Met Lys Lys Ile Arg Glu Ser Tyr Glu

Asn Leu Asn Ala Leu Gly Leu Phe Pro Pro Thr Ser Gly Met Pro Pro

Pro Thr Ser Gly Pro Pro Ser Ala Met Thr Pro Pro Tyr Pro Gln Phe

Asn Asp Ile Ala Ser Met Asn Leu Gln Ala His Leu Ile Pro Gly Leu

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Glu Gln Ser Glu Thr Glu Thr Val His Gln Phe Ile Pro Ala Leu Ser
                                    410
Val Gly Ala Ile Ile Gly Lys Gln Gly Gln His Ile Lys Gln Leu Ser
            420
                                425
Arg Phe Ala Gly Ala Ser Ile Lys Ile Ala Pro Ala Glu Ala Pro Asp
                            440
        435
                                                 445
Ala Lys Val Arg Met Val Ile Ile Thr Gly Pro Pro Glu Ala Gln Phe
    450
                        455
Lys Ala Gln Gly Arg Ile Tyr Gly Lys Ile Lys Glu Glu Asn Phe Val
465
                                        475
                    470
Ser Pro Lys Glu Glu Val Lys Leu Glu Ala His Ile Arg Val Pro Ser
                485
                                    490
Phe Ala Ala Gly Arg Val Ile Gly Lys Gly Gly Lys Thr Val Asn Glu
            500
                                505
                                                     510
Leu Gln Asn Leu Ser Ser Ala Glu Val Val Val Pro Arg Asp Gln Thr
                            520
                                                 525
Pro Asp Glu Asn Asp Gln Val Val Val Lys Ile Thr Gly His Phe Tyr
                        535
                                             540
Ala Cys Gln Val Ala Gln Arg Lys Ile Gln Glu Ile Leu Thr Gln Val
                    550
                                        555
Lys Gln His Gln Gln Gln Lys Ala Leu Gln Ser Gly Pro Pro Gln Ser
                565
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Arg Arg Lys
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<211> 401
<212> DNA
<213> Homo sapiens
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agatccaaac aaatacacat tctgtgtttt agctcagtgt tttctaaaaa aagaaactgc 120
cacacagcaa aaaattgttt actttgttgg acaaaccaaa tcagttctca aaaaatgacc 180
ggtgcttata aaaagttata aatatcgagt agctctaaaa caaaccacct gaccaagagg 240
gaagtgaget tgtgettagt atttacattg gatgeeagtt ttgtaateae tgaettatgt 300
gcaaactggt gcagaaattc tataaactct ttgctgtttt tgatacctgc tttttgtttc 360
attttgtttt gttttgtaaa aatgataaaa cttcagaaaa t
                                                                   401
<210> 178
<211> 561
<212> DNA
<213> Homo sapiens
<400> 178
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geoegetatg ggacaggggt etttggecag aatgagtaee taegetatea ggaggeeetg 120
agtgagctgg ccactgcggt taaagcacga attgggagct ctcagcgaca tcaccagtca 180
gcagccaaag acctaactca gtcccctgag gtctccccaa caaccatcca ggtgacatac 240
ctcccctcca gtcagaaqag taaacgtgcc aagcacttcc ttgaattgaa gagctttaag 300
gataactata acacattgga gagtactctg tgacggagct gaaggactct tgccgtagat 360
taagccagtc agttgcaatg tgcaagacag gctgcttgcc gggccqccct cggaacatct 420
ggcccagcag gcccagactg tatccatcca agttcccgtt gtatccagag ttcttagage 480
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gactattttc ccccagtagc g
<210> 179
<211> 521
<212> DNA
<213> Homo sapiens
<400> 179
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gategageaa tggetteagg acatgggtte tetteteetg tgateattea agtgeteaet 120
qcatqaaqac tqqcttqtct caqtqtttca acctcaccag ggctgtctct tggtccacac 180
ctcqctccct qttaqtqccq tatqacaqcc cccatcaaat gaccttggcc aagtcacggt 240
ttctctqtqq tcaaqqttqq ttqqctqatt qqtqqaaaqt aqqqtqqacc aaagqaqqcc 300
acqtqaqcaq tcaqcaccaq ttctqcacca qcaqcqcctc cqtcctaqtq qqtqttcctq 360
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aggataagtg ggatctacca attgattctg gcaaaacaat ttctaagatt tttttgcttt 480
                                                                521
atgtgggaaa cagatctaaa tctcatttta tgctgtattt t
<210> 180
<211> 417
<212> DNA
<213> Homo sapiens
<400> 180
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tectgggeeg cetggeggee ategtggeta aacaggtact getgggeegg aaggtggtgg 120
tegtaegetg tgaaggeate aacatttetg geaattteta eagaaacaag ttgaagtaee 180
tggctttcct ccgcaagcgg atgaacacca accettcccg aggcccctac cacttccggg 240
ccccagccg catcttctgg cggaccgtgc gaggtatgct gccccacaaa accaagcgag 300
gecaggeege tetggaeegt etcaaggtgt ttgaeggeat eccaeegeee taegaeaaga 360
aaaagcggat ggtggttcct gctgccctca aggtcgtgcg tctgaagcct acaagaa
<210> 181
<211> 283
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 35
<223> n = A, T, C or G
<400> 181
gatttettet aaataggatg taaaaettet tteanattae tetteeteag teetgeetge 60
caagaactca agtgtaactg tgataaaata acctttccca ggtatattgg caggtatgtg 120
atttacattg tttacacttc tatgaccagg ccttaaggga aggtcagttt tttaaaaaaac 240
                                                                 283
caagtagtgt cttcctacct atctccagat acatgtcaaa aaa
<210> 182
<211> 401
<212> DNA
<213> Homo sapiens
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<400> 182
atattettge tgettatgea getgaeattg ttgeeeteee taaageaace aagtageett 60
tatttcccac agtgaaagaa aacgctggcc tatcagttac attacaaaag gcagatttca 120
agaggattga gtaagtagtt ggatggcttt cataaaaaca agaattcaag aagaggattc 180
atgctttaag aaacatttgt tatacattcc tcacaaatta tacctgggat aaaaactatg 240
tagcaggcag tgtgttttcc ttccatgtct ctctgcacta cctgcagtgt gtcctctgag 300
gctgcaagtc tgtcctatct gaattcccag cagaagcact aagaagctcc accctatcac 360
ctagcagata aaactatggg gaaaacttaa atctgtgcat a
                                                                                                                                                 401
<210> 183
<211> 366
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 325
<223> n = A, T, C or G
<400> 183
accepted agentiated accepted a
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tttaaggaca aagatgaagt cactgtaaac taatctgtca ttgtttttac cttccttttc 180
tttttcagtg cagaaattaa aagtaagtat aaagcaccgt gattgggagt gtttttgcgt 240
gtgtcggaat cactggtaaa tgttggctga gaacaatccc tccccttgca cttgtgaaaa 300
cactttgagc gctttaagag attancctga gaaataatta aatatctttt ctcttcaaaa 360
aaaaaa
                                                                                                                                                 366
<210> 184
<211> 370
<212> DNA
<213> Homo sapiens
<400> 184
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tttaataatt gtactgagag aaactgctta cgtacacatt gcagatcaaa tatttggagt 120
taaaatgtta gtctacatag atgggtgatt gtaactttat tgccattaaa agatttcaaa 180
ttgcattcat gcttctgtgt acacataatg aaaaatgggc aaataatgaa gatctctcct 240
tcagtctgct ctgtttaatt ctgctgtctg ctcttctcta atgctgcgtc cctaattgta 300
cacagtttag tgatatctag gagtataaag ttgtcgccca tcaataaaaa tcacaaagtt 360
ggtttaaaaa
                                                                                                                                                370
<210> 185
<211> 107
<212> DNA
<213> Homo sapiens
<400> 185
ctcatattat tttccttttg agaaattgga aactctttct gttgctatta tattaataaa 60
gttggtgttt attttctggt agtcaccttc cccatttaaa aaaaaaa
                                                                                                                                                107
<210> 186
<211> 309
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<212> DNA
<213> Homo sapiens
<400> 186
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agagggccac aggggtggcc gggagttgtc agctgatgcc tgctgagagg caggaattgt 120
gccagtgagt gacagtcatg agggagtgtc tcttcttggg gaggaaagaa ggtagagcct 180
ttctgtctga atgaaaggcc aaggctacag tacagggccc cgccccagcc agggtgttaa 240
tgcccacgta gtggaggcct ctggcagatc ctgcattcca aggtcactgg actgtacgtt 300
tttatggtt
                                                                   309
<210> 187
<211> 477
<212> DNA
<213> Homo sapiens
<400> 187
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tocaaceteg ggccagtgtc ttcaggettt actggggacc tgcgagctgg cctaatgtgg 120
tggcctgcaa gccaggccat ccctgggcgc cacagacgag ctccgagcca ggtcaggctt 180
eggaggeeae aageteagee teaggeeeag geactgattg tggeagaggg geeactaeee 240
aaggtctagc taggcccaag acctagttac ccagacagtg agaagcccct ggaaggcaga 300
aaagttggga gcatggcaga cagggaaggg aaacattttc agggaaaaga catgtatcac 360
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<210> 188
<211> 220
<212> DNA
<213> Homo sapiens
<400> 188
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cagatgttca agaggaagtt gctattgcat tgattttaat atttgtacat aaacactgat 180
ttttttgagc attattttgt atttgttgta ctttaatacc
                                                                   220
<210> 189
<211> 417
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 76, 77
<223> n = A, T, C or G
<400> 189
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ccatcattaa gcatcnnttt caaaattata gccattcatg atttactttt tccagatgac 120
tatcattatt ctagtccttt gaatttgtaa ggggaaaaaa aacaaaaaca aaaacttacg 180
atgcactttt ctccagcaca tcagatttca aattgaaaat taaagacatg ctatggtaat 240
gcacttgcta gtactacaca ctttgtacaa caaaaaacag aggcaagaaa caacggaaag 300
agaaaagcct teettigtig geeettaaae tgagteaaga teigaaatgt agagatgate 360
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<210> 190
<211> 497
<212> DNA
<213> Homo sapiens
<400> 190
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acggtccgca aggatgccta catgttctgg tggctctatt atgccaccaa ctcctgcaag 180
aactteteag aactgeeect ggteatgtgg etteagggeg gteeaggegg ttetageaet 240
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acctggctcc aggctgccag tctcctattt gtggataatc ccgtgggcac tgggttcagt 360
tatgtgaatg gtagtggtgc ctatgccaag gacctggcta tggtggcttc agacatgatg 420
gttctcctga agaccttctt cagttgccac aaagaattcc agacagttcc attctacatt 480
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<210> 191
<211> 175
<212> DNA
<213> Homo sapiens
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gatacccagc attcaataga gaccacacaa taaatatatg tcaaataaaa aaaaa
<210> 192
<211> 526
<212> DNA
<213> Homo sapiens
<400> 192
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attgaagaaa gagaaacttg tcaactcata tccacgttat ctagcaaagt acataagaat 180
ctatcactaa gtaatgtatc cttcagaatg tgttggttta ccagtgacac cccatattca 240
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ttacttaatg tattttggtg tattttcctc aaattaatat tggtgttcaa gactatatct 420
aattoototg atoactitga gaaacaaact titattaaat gtaaggcact titotatgaa 480
ttttaaatat aaaaataaat attgttctga ttattactga aaaaaa
                                                                   526
<210> 193
<211> 553
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 290, 300, 411, 441
<223> n = A, T, C or G
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cagtggtagc agttggactg accattgctg ctgcaggatt tqcaggccqt tacqttttqc 180
aagccatgaa gcatatggag cctcaagtaa aacaagtttt tcaaagccta ccaaaatctg 240
cetteagtgg tggctattat agaggtgggt ttgaacccaa aatgacaaan egggaagcan 300
cattaatact aggtgtaagc cctactgcca ataaagggaa aataagagat gctcatcgac 360
gaattatgct tttaaatcat cctgacaaag gaggatctcc ttatataqca nccaaaatca 420
atgaagctaa agatttacta naaggtcaag ctaaaaaaatg aagtaaatgt atgatgaatt 480
ttaagttcgt attagtttat gtatatgagt actaagtttt tataataaaa tgcctcagag 540
ctacaatttt aaa
                                                                   553
<210> 194
<211> 320
<212> DNA
<213> Homo sapiens
<400> 194
cccttcccaa tccatcagta aagaccccat ctgccttgtc catgccgttt cccaacaggg 60
atgtcacttg atatgagaat ctcaaatctc aatgccttat aagcattcct tcctqtqtcc 120
attaagactc tgataattgt ctcccctcca taggaatttc tcccaggaaa gaaatatatc 180
cccatctccg tttcatatca gaactaccgt ccccgatatt cccttcaqaq aqattaaaqa 240
ccagaaaaaa gtgagcctct tcatctgcac ctgtaatagt ttcagttcct attttcttcc 300
attgacccat atttatacct
                                                                   320
<210> 195
<211> 320
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 203, 218
<223> n = A, T, C or G
<400> 195
aagcatgacc tggggaaatg gtcagacctt gtattgtgtt tttggccttg aaagtagcaa 60
gtgaccagaa tctgccatgg caacaggctt taaaaaaagac ccttaaaaaag acactgtctc 120
aactgtggtg ttagcaccag ccagctctct gtacatttqc tagcttqtag ttttctaaga 180
ctgagtaaac ttcttatttt tanaaagggg aggctggntt gtaactttcc ttgtacttaa 240
ttgggtaaaa gtcttttcca caaaccacca tctattttgt gaactttgtt agtcatcttt 300
tatttggtaa attatgaact
                                                                   320
<210> 196
<211> 357
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 36
<223> n = A, T, C or G
<400> 196
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THE THE ACT OF THE ACT
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atataaaata atacqaaact ttaaaaaagca ttqqantqtc aqtatqttqa atcaqtaqtt 60
tcactttaac tgtaaacaat ttcttaggac accatttggg ctagtttctg tgtaagtgta 120
aatactacaa aaacttattt atactgttct tatgtcattt gttatattca tagatttata 180
tgatgatatg acatetgget aaaaagaaat tattgcaaaa etaaceaeta tgtaettttt 240
tataaatact gtatggacaa aaaatggcat titttatatt aaattgttta gctctggcaa 300
aaaaaaaaa ttttaagagc tggtactaat aaaggattat tatgactgtt aaaaaaa
<210> 197
<211> 565
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 27
<223> n = A, T, C or G
<400> 197
teagetgagt accateagga tatttaneee tttaagtget gttttgggag tagaaaacta 60
aagcaacaat actteetett gacagetttg attggaatgg ggttattaga teatteacet 120
tggtcctaca ctttttagga tgcttggtga acataacacc acttataatg aacatccctg 180
gttcctatat tttgggctat gtgggtagga attgttactt gttactgcag cagcagccct 240
agaaagtaag cccagggctt cagatctaag ttagtccaaa agctaaatga tttaaagtca 300
agttgtaatg ctaggcataa gcactctata atacattaaa ttataggccg agcaattagg 360
gaatgtttct gaaacattaa acttgtattt atgtcactaa aattctaaca caaacttaaa 420
aaatgtgtct catacatatg ctgtactagg cttcatcatg catttctaaa tttgtgtatg 480
atttgaatat atgaaagaat ttatacaaga gtgttattta aaattattaa aaataaatgt 540
                                                                 565
atataatttq tacctattqt aaaaa
<210> 198
<211> 484
<212> DNA
<213> Homo sapiens
<400> 198
tatgtaagta ttggtgtctg ctttaaaaaa ggagacccag acttcacctg tcctttttaa 60
acatttgaga acagtgttac tetgagcagt tgggccaect teacettate egacagetga 120
tgggcgcagc agcaggtggc aggggtgtgg cttgaggtgg gtggcagcgt ctqqtcctcc 240
tetetggtge tttetgagag ggtetetaaa geagagtgtg gttggeetgg gggaaggeag 300
ageaegtatt teteceetet agtacetetg cattigtgag tgttecetet ggetttetga 360
agggcagcag actettgagt atactgcaga ggacatgett tatcagtagg teetgaggge 420
tccaggggct caactgacca agtaacacag aagttggggt atgtggccta tttgggtcgg 480
aaac
                                                                 484
<210> 199
<211> 429
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 77, 88, 134, 151, 189, 227, 274, 319
<223> n = A, T, C or G
```

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<400> 199
gcttatgttt tttgttttaa cttttgtttt ttaacattta gaatattaca ttttgtatta 60
tacagtacct ttctcanaca ttttgtanaa ttcatttcgg cagctcacta ggattttgct 120
gaacattaaa aagngtgata gcgatattag ngccaatcaa atggaaaaaa ggtagtctta 180
ataaacaana cacaacgttt ttatacaaca tactttaaaa tattaanaaa actccttaat 240
attgtttcct attaagtatt attctttggg caanattttc tgatgctttt gattttctct 300
caatttagca tttgctttng gtttttttct ctatttagca ttctgttaag gcacaaaaac 360
tatgtactgt atgggaaatg ttgtaaatat taccttttcc acattttaaa cagacaactt 420
                                                                   429
tgaatccaa
<210> 200
<211> 279
<212> DNA
<213> Homo sapiens
<400> 200
gcttttttga ggaattacag ggaagctcct ggaattgtac atggatatct ttatccctag 60
ggggaaatca aggagctggg cacccctaat tetttatgga agtgtttaaa actattttaa 120
ttttattaca agtattacta gagtagtggt tctactctaa gatttcaaaa gtgcatttaa 180
aatcatacat gttcccgcct gcaaatatat tgttattttg gtggagaaaa aaatagtata 240
ttctacataa aaaattaaag atattaacta agaaaaaaa
                                                                   279
<210> 201
<211> 569
<212> DNA
<213> Homo sapiens
<400> 201
taggtcagta tttttagaaa ctcttaatag ctcatactct tgataccaaa agcagccctg 60
attqttaaaq cacacacctq cacaaqaaqc aqtqatqqtt gcatttacat ttcctgggtg 120
cacaaaaaaa aattotcaaa aagcaaggac ttacgetttt tgcaaagcct ttgagaagtt 180
actggatcat aggaagctta taacaagaat ggaagattct taaataactc actttctttg 240
gtatccagta acagtagatg ttcaaaatat gtagctgatt aataccagca ttgtgaacgc 300
tgtacaacct tgtggttatt actaagcaag ttactactag cttctgaaaa gtagcttcat 360
aattaatgtt atttatacac tgccttccat gacttttact ttgccctaag ctaatctcca 420
aaatetqaaa tqctactcca atateaqaaa aaaaqqqqqa qqtqqaatta tattteetgt 480
gattttaaga gtacagagaa tcatgcacat ctctgattag ttcatatatg tctagtgtgt 540
aataaaagtc aaagatgaac tctcaaaaa
                                                                   569
<210> 202
<211> 501
<212> DNA
<213> Homo sapiens
<400> 202
attaataggc ttaataattg ttggcaagga tccttttgct ttctttggca tgcaagctcc 60
tagcatctgg cagtggggcc aagaaaataa ggtttatgca tgtatgatgg ttttcttctt 120
gagcaacatg attgagaacc agtgtatgtc aacaggtgca tttgagataa ctttaaatga 180
tgtacctgtg tggtctaagc tggaatctgg tcaccttcca tccatgcaac aacttgttca 240
aattettgae aatgaaatga ageteaatgt geatatggat teaateeeac accategate 300
atagcaccac ctatcagcac tgaaaactct tttgcattaa gggatcattg caagagcagc 360
qtgactgaca ttatgaaggc ctgtactgaa gacagcaagc tgttagtaca gaccagatgc 420
tttcttggca ggctcgttgt acctcttgga aaacctcaat gcaagatagt gtttcagtgc 480
```

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501
tggcatattt tggaattctg c
<210> 203
<211> 261
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 36, 96
<223> n = A, T, C or G
<400> 203
gacaagetee tggtettgag atgtettete gttaangaga tgggeetttt ggaggtaaag 60
gataaaatga atgagttetg teatgattea etattntata aettgeatga eetttaetgt 120
gttagetett tgaatgttet tgaaatttta gaetttettt gtaaacaaat gatatgteet 180
tatcattgta taaaagctgt tatgtgcaac agtgtggaga ttccttgtct gatttaataa 240
aatacttaaa cactgaaaaa a
<210> 204
<211> 421
<212> DNA
<213> Homo sapiens
<400> 204
agcatctttt ctacaacgtt aaaattgcag aagtagctta tcattaaaaa acaacaacaa 60
caacaataac aataaatcct aagtgtaaat cagttattct accccctacc aaggatatca 120
gcctgttttt tccctttttt ctcctgggaa taattgtggg cttcttccca aatttctaca 180
geotetttee tetteteatg ettgagette eetgtttgea egeatgegtg tgeaggaetg 240
gettgtgtge ttggactegg etceaggtgg aageatgett teeettgtta etgttggaga 300
aactcaaacc ttcaagccct aggtgtagcc attttgtcaa gtcatcaact gtatttttgt 360
actggcatta acaaaaaaag aagataaaat attgtaccat taaactttaa taaaacttta 420
                                                                    421
<210> 205
<211> 460
<212> DNA
<213> Homo sapiens
<400> 205
 tactctcaca atgaaggacc tggaatgaaa aatctgtgtc taaacaagtc ctctttagat 60
 tttagtgcaa atccagagcc agcgtcggtt gcctcgagta attctttcat gggtaccttt 120
 ggaaaagete teaggagaee teacetagat geetatteaa getttggaea geeateagat 180
 tgtcagccaa gagcctttta tttgaaagct cattcttccc cagacttgga ctctgggtca 240
 gaggaagatg ggaaagaaag gacagatttt caggaagaaa atcacatttg tacctttaaa 300
 cagactttag aaaactacag gactccaaat tttcagtctt atgacttgga cacatagact 360
 gaatgagacc aaaggaaaag cttaacatac tacctcaagg tgaactttta tttaaaagag 420
                                                                    460
 agagaatett atgtttttta aatggagtta tgaattttaa
 <210> 206
 <211> 481
 <212> DNA
 <213> Homo sapiens
```

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<400> 206
tgtggtggaa ttcgggacgc ccccagaccc tgactttttc ctgcgtgggc cgtctcctcc 60
tgcggaagca gtgacetetg acceetggtg acettegett tgagtgeett ttgaacgetg 120
gtcccgcggg acttggtttt ctcaagctct gtctgtccaa agacgctccg gtcgaggtcc 180
egectgeeet gggtggatae ttgaaceeea gaegeeeete tgtgetgetg tgteeggagg 240
eggeetteec atetgeetge ecaeceggag etettteege eggegeaggg teecaageee 300
acctecegee eteagteetg eggtgtgegt etgggeaegt eetgeacaca caatgeaagt 360
cetggeetee gegeegeee geceaegega geegtaceeg eegeeaacte tgttatttat 420
ggtgtgaccc cctggaggtg ccctcggccc accggggcta tttattgttt aatttatttg 480
                                                                   481
t.
<210> 207
<211> 605
<212> DNA
<213> Homo sapiens
<400> 207
accetttttg gatteaggge teeteacaat taaaatgagt gtaatgaaac aaggtgaaaa 60
tatagaagca tccctttgta tactgttttg ctacttacag tgtacttggc attgctttat 120
ctcactggat teteaeggta ggatttetga gatettaate taageteeaa agttgtetae 180
ttttttgatc ctagggtgct ccttttgttt tacagagcag ggtcacttga tttgctagct 240
ggtggcagaa ttggcaccat tacccaggtc tgactgacca ccagtcagag gcactttatt 300
tgtatcatga aatgatttga aatcattgta aagcagcgaa gtctgataat gaatgccagc 360
tttccttgtg ctttgataac aaagactcca aatattctgg agaacctgga taaaagtttg 420
aagggctaga ttgggatttg aagacaaaat tgtaggaaat cttacatttt tgcaataaca 480
aacattaatg aaagcaaaac attataaaag taattttaat tcaccacata cttatcaatt 540
tettgatget tecaaatgae atetaceaga tatggttttg tggacatett tttetgttta 600
cataa
                                                                   605
<210> 208
<211> 655
<212> DNA
<213> Homo sapiens
<400> 208
ggcgttgttc tggattcccg tcgtaactta aagggaaact ttcacaatgt ccggagccct 60
tgatgtcctg caaatgaagg aggaggatgt ccttaagttc cttgcagcag gaacccactt 120
aggtggcacc aatcttgact tccagatgga acagtacatc tataaaagga aaagtgatgg 180
catctatate ataaatetea agaggaeetg ggagaagett etgetggeag etegtgeaat 240
tgttgccatt gaaaaccctg ctgatgtcag tgttatatcc tccaggaata ctggccagag 300
ggctgtgctg aagtttgctg ctgccactgg agccactcca attgctggcc gcttcactcc 360
tggaaccttc actaaccaga tccaggcagc cttccgggag ccacggcttc ttgtggttac 420
tgaccccagg gctgaccacc agcctctcac ggaggcatct tatgttaacc tacctaccat 480
tgcgctgtgt aacacagatt ctcctctgcg ctatgtggac attgccatcc catgcaacaa 540
caagggaget cacteagtgg gtttgatgtg gtggatgetg getegggaag ttetgegeat 600
gcgtggcacc atttcccgtg aacacccatg ggaggtcatg cctgatctgt acttc
                                                                   655
<210> 209
<211> 621
<212> DNA
<213> Homo sapiens
<400> 209
catttagaac atggttatca tocaagacta ctotaccotg caacattgaa ctoccaagag 60
```

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caaatccaca ttcctcttga gttctgcagc ttctgtgtaa atagggcagc tgtcgtctat 120
gccgtagaat cacatgatct gaggaccatt catggaagct gctaaatagc ctagtctggg 180
gagtetteca taaagttttg catggageaa acaaacagga ttaaactagg tttggtteet 240
teagecetet aaaageatag ggettageet geaggettee ttgggettte tetgtgtgtg 300
tagttttgta aacactatag catctgttaa gatccagtgt ccatggaaac cttcccacat 360
geogtgaete tggaetatat eagtttttgg aaageagggt teetetgeet getaacaage 420
ccacgtggac cagtctgaat gtctttcctt tacacctatg tttttaaata gtcaaacttc 480
aagaaacaat ctaaacaagt ttctgttgca tatgtgtttg tgaacttgta tttgtattta 540
gtaggettet atattgeatt taacttgttt ttgtaactee tgattettee tttteggata 600
                                                                 621
ctattgatga ataaagaaat t
<210> 210
<211> 533
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 20, 21, 61
<223> n = A, T, C or G
<400> 210
cgccttgggg agccggcggn ngagtccggg acgtggagac ccggggtccc ggcagccggg 60
nggccegegg geceagggtg gggatgeace geegeggggt gggagetgge gecategeca 120
agaagaaact tgcagaggcc aagtataagg agcgagggac ggtcttggct gaggaccagc 180
tageceagat gteaaageag ttggaeatgt teaagaeeaa eetggaggaa tttgeeagea 240
aacacaaqca qqaqatccqq aaqaatcctq aqttccqtqt qcaqttccag gacatgtgtg 300
caaccattgg cgtggatccg ctggcctctg gaaaaggatt ttggtctgag atgctgggcg 360
tgggggactt ctattacgaa ctaggtgtcc aaattatcga agtgtgcctg gcgctgaagc 420
gcaagttcgc ccaggatgtc agtcaagatg acctgatcag agccatcaag aaa
<210> 211
<211> 451
<212> DNA
<213> Homo sapiens
<400> 211
ttagcttgag ccgagaacga ggcgagaaag ctggagaccg aggagaccgc ctagagcgga 60
gtgaacgggg aggggaccgt ggggaccggc ttgatcgtgc gcggacacct gctaccaagc 120
ggagcttcag caaggaagtg gaggagcgga gtagagaacg gccctcccag cctgaggggc 180
tgcgcaaggc agctagcete acggaggate gggaccgtgg gcgggatgce gtgaagcgag 240
aagctgccct accccagtg agccccctga aggcggctct ctctgaggag gagttagaga 300
agaaatccaa ggctatcatt gaggaatatc tccatctcaa tgacatgaaa gaggcagtcc 360
agtgcgtgca ggagctggcc tcaccctcct tgctcttcat ctttgtacgg catggtgtcg 420
                                                                 451
agtctacgct ggagcgcagt gccattgctc g
<210> 212
<211> 471
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
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<222> 54
<223> n = A, T, C or G
<400> 212
gtgattattc ttgatcaggg agaagatcat ttagatttgt tttgcattcc ttanaatgga 60
gggcaacatt ccacagctgc cctggctgtg atgagtgtcc ttgcaggggc cggagtagga 120
gcactggggt gggggggaa ttggggttac tcgatgtaag ggattccttg ttgttgtgtt 180
gagatecagt geagttgtga tttetgtgga teccagettg gttecaggaa ttttgtgtga 240
ttggcttaaa tccaqttttc aatcttcgac agctgggctg gaacgtgaac tcagtagctg 300
aacctgtctg acccggtcac gttcttggat cctcagaact ctttgctctt gtcggggtgg 360
gggtgggaac tcacgtgggg agcggtggct gagaaaatgt aaggattctg gaatacatat 420
tocatgggae ttteetteee teteetgett cetetttee tgeteectaa e
                                                                   471
<210> 213
<211> 511
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 27, 63, 337, 442
<223> n = A, T, C or G
<400> 213
ctaattagaa acttgctgta ctttttnttt tcttttaggg gtcaaggacc ctctttatag 60
ctnccatttg cctacaataa attattgcag cagtttgcaa tactaaaata ttttttatag 120
actttatatt tttccttttg ataaagggat gctgcatagt agagttggtg taattaaact 180
ateteageeg titeeetget tieeettetg etecatatge eteatigtee tieeagggag 240
ctcttttaat cttaaagttc tacatttcat gctcttagtc aaattctgtt acctttttaa 300
taactettee cactgeatat ttecatettg aattggnggt tetaaattet gaaactgtag 360
ttgagataca gctatttaat atttctggga gatgtgcatc cctcttcttt gtggttgccc 420
aaggttgttt tgcgtaactg anactccttg atatgcttca gagaatttag gcaaacactg 480
gccatggccg tgggagtact gggagtaaaa t
                                                                   511
<210> 214
<211> 521
<212> DNA
<213> Homo sapiens
<400> 214
agcattgcca aataatccct aattttccac taaaaatata atgaaatgat gttaagcttt 60
ttgaaaagtt taggttaaac ctactgttgt tagattaatg tatttgttgc ttccctttat 120
ctggaatgtg gcattagctt ttttatttta accetettta attettatte aattecatga 180
cttaaggttg gagagctaaa cactgggatt tttggataac agactgacag ttttgcataa 240
ttataategg cattgtacat agaaaggata tggctacett ttgttaaate tgcactttet 300
aaatatcaaa aaagggaaat gaagtataaa tcaatttttg tataatctgt ttgaaacatg 360
agttttattt gcttaatatt agggetttge ceettttetg taagtetett gggateetgt 420
gtagaagetg tteteattaa acaccaaaca gttaagteca ttetetggta etagetacaa 480
attcggtttc atattctact taacaattta aataaactga a
                                                                   521
<210> 215
<211> 381
<212> DNA
<213> Homo sapiens
```

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<220>
<221> misc feature
<222> 17, 20, 60, 61, 365
<223> n = A, T, C or G
<400> 215
gageggagag eggacength agageeetga geageeecae egeegeegee ggeetagtth 60
ncatcacacc cogggaggag cogcagctgc cgcagcoggc cocagtcacc atcaccgcaa 120
ccatgagcag cgaggccgag acccagcage cgcccgccgc cccccccgcc gcccccgccc 180
teagegeege egacaeeaag eeeggeacta egggeagegg egeagggage ggtggeeegg 240
geggeeteae ateggeggeg eetgeeggeg gggacaagaa ggteategea aegaaggttt 300
tgggaacagt aaaatggttc aatgtaagga acggatatgg tttcatcaac aggaatgaca 360
ccaangaaga tgtatttgta c
                                                                   381
<210> 216
<211> 425
<212> DNA
<213> Homo sapiens
<400> 216
ttactaacta ggtcattcaa ggaagtcaag ttaacttaaa catgtcacct aaatgcactt 60
gatggtgttg aaatgtccac cttcttaaat ttttaagatg aacttagttc taaagaagat 120
aacaggccaa teetgaaggt acteeetgtt tgetgeagaa tgteagatat tttggatgtt 180
gcataagagt cetatttgcc ccagttaatt caacttttgt ctgcctgttt tgtggactgg 240
ctggctctgt tagaactctg tccaaaaagt gcatggaata taacttgtaa agcttcccac 300
aattgacaat atatatgcat gtgtttaaac caaatccaga aagcttaaac aatagagctg 360
cataatagta tttattaaag aatcacaact gtaaacatga gaataactta aggattctag 420
                                                                   425
<210> 217
<211> 181
<212> DNA
<213> Homo sapiens
<400> 217
gagaaaccaa atgataggtt gtagagcctg atgactccaa acaaagccat cacccgcatt 60
cttcctcctt cttctggtgc tacagctcca agggcccttc accttcatgt ctgaaatgga 120
actttggctt tttcagtgga agaatatgtt gaaggtttca ttttgttcta gaaaaaaaa 180
                                                                   181
<210> 218
<211> 405
<212> DNA
<213> Homo sapiens
<400> 218
caggeettee agtteactga caaacatggg gaagtgtgee cagetggetg gaaacetgge 60
agtgatacca tcaagcctga tgtccaaaag agcaaagaat atttctccaa gcagaagtga 120
gegetggget gttttagtge eaggetgegg tgggeageea tgagaacaaa acetettetg 180
tatttttttt ttccattagt aaaacacaag acttcagatt cagccgaatt gtggtgtctt 240
acaaggcagg cettteetae agggggtgga gagaccagee tttetteett tggtaggaat 300
ggcctgagtt ggcgttgtgg gcaggctact ggtttgtatg atgtattagt agagcaaccc 360
attaatcttt tgtagtttgt attaaacttg aactgagaaa aaaaa
                                                                   405
```

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<210> 219
<211> 216
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 207, 210
<223> n = A, T, C or G
<400> 219
actccaagag ttagggcagc agagtggagc gatttagaaa gaacatttta aaacaatcag 60
ttaatttacc atqtaaaatt qctqtaaatq ataatqtqta caqattttct qttcaaatat 120
tcaattgtaa acttcttgtt aagactgtta cgtttctatt gcttttgtat gggatattgc 180
aaaaataaaa aggaaagaac cctcttnaan aaaaaa
                                                                   216
<210> 220
<211> 380
<212> DNA
<213> Homo sapiens
<400> 220
cttacaaatt gcccccatgt gtaggggaca cagaaccctt tgagaaaact tagatttttg 60
tetgtacaaa gtetttgeet tttteettet teattttttt ceagtacatt aaatttgtea 120
atttcatctt tgagggaaac tgattagatg ggttgtgttt gtgttctgat ggagaaaaca 180
gcaccccaag gactcagaag atgattttaa cagttcagaa cagatgtgtg caatattggt 240
gcatgtaata atgttgagtg gcagtcaaaa gtcatgattt ttatcttagt tcttcattac 300
tgcattgaaa aggaaaacct gtctgagaaa atgcctgaca gtttaattta aaactatggt 360
gtaagtcttt gacaaaaaa
                                                                    380
<210> 221
<211> 398
<212> DNA
<213> Homo sapiens
<400> 221
ggttagtaag ctgtcgactt tgtaaaaaag ttaaaaatga aaaaaaaagg aaaaatgaat 60
tgtatattta atgaatgaac atgtacaatt tgccactggg aggaggttcc tttttgttgg 120
gtgagtctgc aagtgaattt cactgatgtt gatattcatt gtgtgtagtt ttatttcggt 180
eccageceeg titteetitta tittiggaget aatgecaget gegigtetag tittigagige 240
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<221> misc feature
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<223> n = A, T, C or G
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gtgaagattt caaaacctga gagcactttt tctttgttta gaattatgag aaaggcacta 180
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<213> Homo sapiens
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gctggatgaa cttaaaaaaa
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<211> 385
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tetecaacae cageaageee taaccaggge eeteeteeae aagtteeagt ateteetgga 180
ccaccaaagg acagttctgc ccctggtgga cccccagaaa ggactgttac tccagcccta 240
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<213> Homo sapiens
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Arg Pro Ser Ala Tyr Met Arg Glu His Asn Gln Leu Asn Gly Trp Ser
                            40
Ser Asp Glu Asn Asp Trp Asn Glu Lys Leu Tyr Pro Val Trp Lys Arg
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Gly Asp Met Arg Trp Lys Asn Ser Trp Lys Gly Gly Arg Val Gln Ala
                    70
                                         75
Val Leu Thr Ser Asp Ser Pro Ala Leu Val Gly Ser Asn Ile Thr Phe
                                     90
Ala Val Asn Leu Ile Phe Pro Arg Cys Gln Lys Glu Asp Ala Asn Gly
            100
                                 105
                                                     110
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Asp	Pro 130	Tyr	Val	Tyr	Asn	Trp 135	Thr	Ala	Trp	Ser	Glu 140	Asp	Ser	Asp	Gly
Glu 145	Asn	Gly	Thr	Gly	Gln 150	Ser	His	His	Asn	Val 155	Phe	Pro	Asp	Gly	Lys 160
Pro	Phe	Pro	His	His 165	Pro	Gly	Trp	Arg	Arg 170	Trp	Asn	Phe	Ile	Tyr 175	Val
Phe	His	Thr	Leu 180	Gly	Gln	Tyr	Phe	Gln 185	Lys	Leu	Gly	Arg	Cys 190	Ser	Val
Arg	Val	Ser 195	Val	Asn	Thr	Ala	Asn 200	Val	Thr	Leu	Gly	Pro 205	Gln	Leu	Met
Glu	Val 210	Thr	Val	Tyr	Arg	Arg 215	His	Gly	Arg	Ala	Tyr 220	Val	Pro	Ile	Ala
Gln 225	Val	Lys	Asp	Val	Tyr 230	Val	Val	Thr	Asp	Gln 235	Ile	Pro	Val	Phe	Val 240
Thr	Met	Phe	Gln	Lys 245	Asn	Asp	Arg	Asn	Ser 250	Ser	Asp	Glu	Thr	Phe 255	Leu
Lys	Asp	Leu	Pro 260	Ile	Met	Phe	Asp	Val 265	Leu	Ile	His	Asp	Pro 270	Ser	His
Phe	Leu	Asn 275	Tyr	Ser	Thr	Ile	Asn 280	Tyr	Lys	Trp	Ser	Phe 285	Gly	Asp	Asn
Thr	Gly 290	Leu	Phe	Val	Ser	Thr 295	Asn	His	Thr	Val	Asn 300	His	Thr	Tyr	Val
Leu 305	Asn	Gly	Thr	Phe	Ser 310	Leu	Asn	Leu	Thr	Val 315	Lys	Ala	Ala	Ala	Pro 320
Gly	Pro	Cys	Pro	Pro 325	Pro	Pro	Pro	Pro	Pro 330	Arg	Pro	Ser	Lys	Pro 335	Thr
Pro	Ser	Leu	Gly 340	Pro	Ala	Gly	Asp	Asn 345	Pro	Leu	Glu	Leu	Ser 350	Arg	Ile
Pro	Asp	Glu 355	Asn	Cys	Gln	Ile	Asn 360	Arg	Tyr	Gly	His	Phe 365	Gln	Ala	Thr
Ile	Thr 370	Ile	Val	Glu	Gly	Ile 375	Leu	Glu	Val	Asn	Ile 380	Ile	Gln	Met	Thr
385			Met		390					395				=	400
			Cys	405					410					415	
Ser	Asp	Pro	Thr 420	Cys	Glu	Ile	Thr	Gln 425	Asn	Thr	Val	Cys	Ser 430	Pro	Val
Asp	Val	Asp 435	Glu	Met	Cys	Leu	Leu 440	Thr	Val	Arg	Arg	Thr 445	Phe	Asn	Gly
	450		Tyr			455					460				
Ala 465	Leu	Thr	Ser	Thr	Leu 470	Ile	Ser	Val	Pro	Asp 475	Arg	Asp	Pro	Ala	Ser 480
Pro	Leu	Arg	Met	Ala 485	Asn	Ser	Ala	Leu	Ile 490	Ser	Val	Gly	Cys	Leu 495	Ala
			Thr 500					505					510	_	
-	Asn	515	Ile				520					525		_	_
Leu	Ser 530	Val	Phe	Leu	Asn	Arg 535	Ala	Lys	Ala	Val	Phe 540	Phe	Pro	Gly	Asn

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Gln Glu Lys Asp Pro Leu Leu Lys Asn Gln Glu Phe Lys Gly Val Ser
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     <400> 227
     Phe Leu Leu Asn Asp Asn Leu Thr Ala
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      1
i . j
i g
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<211> 9
ź
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175
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151
S. H. Sref. stoll '
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                   5
ē = £
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     <211> 10
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Phe Ser Phe Ala
All Hand Tour I
                  20
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     <211> 21
1,1
     <212> PRT
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£
1 TA
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10
Bull Hack
     Asn His Ser Pro Ser
              20
i =1:
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Phe Ile Pro Pro Asn
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Arg Asn Pro Gln Gln Ala Gly Ile Arg Glu Ile Phe Thr Phe Ser Pro
Gln Ile Ser Thr
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<210> 240
<211> 21
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Series and American
                                  10
     Val Leu Gly Val
ir with this i
                  20
1.0
     <210> 242
1.
     <211> 20
113
     <212> PRT
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ë
1 1 1
     <400> 242
17
     Gly Ser His Ala Met Tyr Val Pro Gly Tyr Thr Ala Asn Gly Asn Ile
Ting.
     1
                                         10
ij,
     Gln Met Asn Ala
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ä=8:
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                                          10
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His Phe Pro His
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                                        10
     Gln Ala Leu Lys
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ığı
<400> 246
1.0
     Asn Leu Thr Phe Arg Thr Ala Ser Leu Trp Ile Pro Gly Thr Ala Lys
1.1
     1
                                        10
17
     Pro Gly His Trp
                 20
E
note that
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Hum
Hum
     <211> 20
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125
     <213> Homo sapiens
i sk
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     Phe Tyr Pro Ile
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                               25
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                           40
Ala Ala Val Ser Ser Ile Phe Asn Ser Pro Glu Glu Phe Leu Gly Lys
                       55
Ala Val Gly Leu Ser Ala Glu Ala Leu Thr Ile Gln Gln Tyr Ala Asp
                   70
                                       75
Val Leu Ser Lys Ala Leu Gly Lys Glu Val Arg Asp Ala Lys Ile Thr
                                   90
Pro Glu Ala Phe Glu Lys Leu Gly Phe Pro Ala Ala Lys Glu Ile Ala
```

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100

```
Asn Met Cys Arg Phe Tyr Glu Met Lys Pro Asp Arg Asp Val Asn Leu
        115
                            120
                                                125
Thr His Gln Leu Asn Pro Lys Val Lys Ser Phe Ser Gln Phe Ile Ser
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8031

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<210> 255
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
\langle 222 \rangle 9, 6\overline{7}, 247, 275, 277, 397
<223> n = A, T, C or G
<400> 255
gtggccagng actagaaggc gaggcgccgc gggaccatgg cggcggcggc ggacgagcgg 60
agtccanagg acggagaaga cgaggaagag gaggagcagt tqqttctqqt qqaattatca 120
ggaattattg attcagactt cctctcaaaa tgtgaaaata aatgcaaggt tttgggcatt 180
gacactgaga ggcccattct gcaagtggac agctgtgtct ttgctgggga gtatgaagac 240
actetangga eetgtgttat atttgaagaa aatgntnaae atgetgatae agaaggeaat 300
aataaaacag tgctaaaata taaatgccat acaatgaaga agctcagcat gacaagaact 360
ctcctgacag agaagaagga aggagaagaa aacatangtg g
<210> 256
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 7, 37, 51, 79, 96, 98, 103, 104, 107, 116, 167, 181, 183,
194, 206, 276, 303, 307, 308, 310, 323, 332, 341, 353, 374,
376
<223> n = A, T, C or G
<400> 256
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gggccggggt cgcggccgng gacggggccg gggccnangc cgnnganctc gcggangcaa 120
ggccgaggat aaggagtgga tgcccgtcac caacttgggc cgcttgncca aggacatgaa 180
nancaagccc ctgnaggaga tctatntctt cttccctqcc ccattaagga atcaagagat 240
catttgattt cttcctgggg gcctctctca aggatnaggt ttttgaagat tatgccagtg 300
canaaannan acccegttge cengteeate theacceaae netteeaagg genatttttg 360
tttaggcctc attncngggg ggaaccttaa cccaatttgg g
                                                                     401
<210> 257
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 382, 387
\langle 223 \rangle n = A, T, C or G
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<400> 257
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ctctcagccc tgaggtatac agaatcattt gcctcagact gctgttggat tttaaaattt 120
ttaaaatatc tgctaagtaa tttgctatgt cttctcccac actatcaata tgcctqcttc 180
taacaggete eccaetttet tttaatgtge tgttatgage tttggacatg agataacegt 240
gcctgttcag agtgtctaca gtaagagctg gacaaactct ggagggacac agtctttgag 300
acagetettt tggttgettt ceaettttet gaaaggttea eagtaacett etagataata 360
gaaactccca gttaaagcct angctancaa tttttttag t
                                                                   401
<210> 258
<211> 401
<212> DNA
<213> Homo sapiens
<400> 258
ggagcgctag gtcggtgtac gaccgagatt agggtgcgtg ccagctccgg gaggccgcgg 60
tgaggggccg ggcccaagct gccgacccga gccgatcgtc agggtcgcca gcgcctcagc 120
tctgtggagg agcagcagta gtcggagggt gcaggatatt agaaatggct actccccaqt 180
caattttcat ctttgcaatc tgcattttaa tgataacaga attaattctg gcctcaaaaa 240
gctactatga tatcttaggt gtgccaaaat cggcatcaga gcgccaaatc aagaaggcct 300
ttcacaagtt ggccatgaag taccaccctg acaaaaataa gacccagatg ctgaagcaaa 360
attcagagag attgcagaag catatgaaac actctcagat g
                                                                   401
<210> 259
<211> 401
<212> DNA
<213> Homo sapiens
<400> 259
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ctccagaata ttgtgggttt gatcatcaat gcagtcatgt taggctgcat tttcatgaaa 120
acageteagg eteacagaag ggeagaaact ttgattttea geegeeatge tgtgattgee 180
gtccgaaatg gcaagctgtg cttcatgttc cgagtgggtg acctgaggaa aagcatgatc 240
attagtgcct ctgtgcgcat ccaggtggtc aagaaaacaa ctacacctga aggggaggtg 300
gttcctattc accaactgga cattcctgtt gataacccaa tcgagagcaa taacattttt 360
ctggtggccc ctttgatcat ctgccacqtq attgacaaqc q
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<210> 260
<211> 363
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 7, 9, 19, 41, 63, 73, 106, 111, 113, 116, 119, 156, 158,
162, 187, 247, 288, 289, 290, 292, 298, 299, 300, 340
<223> n = A, T, C or G
<400> 260
aggaganang gagggggana tgaataggga tggagaggga natagtggat gagcagggca 60
canggagagg aancagaaag gagaggcaag acagggagac acacancaca nangangana 120
caggtggggg ctggggtggg gcatggagag cctttnangt cncccaggcc accetgctct 180
cgctggnctg ttgaaaccca ctccatggct tcctgccact gcagttgggc ccagggctgg 240
cttattnctg gaatgcaagt ggctgtggct tggagcctcc cctctggnnn anggaaannn 300
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aca
<210> 261
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 114, 152
<223> n = A, T, C or G
<400> 261
eggeteteeg eegeteteee ggggtttegg ggeacttggg teecacagte tggteetget 60
tcaccttccc ctgacctgag tagtcgccat ggcacaggtt ctcagaggca ctgngactga 120
cttccctgga tttgatgagc gggctgatgc anaaactctt cggaaggcta tgaaaggctt 180
gggcacagat gaggagagca tectgactet gttgacatee egaagtaatg eteagegeea 240
ggaaatetet geagetttta agaetetgtt tggeagggat ettetggatg acetgaaate 300
agaactaact ggaaaatttg aaaaattaat tgtggctctg atgaaaccct ctcggcttta 360
tgatgcttat gaactgaaac atgccttgaa gggagctgga a
                                                                   401
<210> 262
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 7, 26, 258, 305, 358, 373, 374, 378
<223> n = A, T, C or G
<400> 262
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tttttaaata ctgtaaagtg acatatagtt ataagatata tttctgtaca gtagagaaag 120
agtttataac atgaagaata ttgtaccatt atacattttc attctcgatc tcataagaaa 180
ttcaaaagaa taatgataga ggtgaaaata tgtttacttt ctctaaatca agcctagttg 240
tcaactcaaa aattatgntg catagtttta ttttgaattt aggttttggg actacttttt 300
tocanottoa atgagaaaat aaaatotaca actoaggagt tactacagaa gttotaanta 360
tttttttgct aannagcnaa aaatataaac atatgaaaat g
                                                                   401
<210> 263
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 232, 290, 304, 326, 383
<223> n = A, T, C or G
<400> 263
ctgtccgacc aagagaggcc ggccgagccc gaggcttggg cttttgcttt ctggcggagg 60
gatetgegge ggtttaggag geggegetga teetgggagg aagaggeage taeggeggeg 120
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geggeggtgg eggetaggge ggeggegaat aaaggggeeg eegeegggtg atgeggtgae 180
cactgeggea ggeecaggag etgagtggge eeeggeeete ageeegteee gneggaeeeg 240
ctttcctcaa ctctccatct tctcctgccg accgagatcg ccgaggcggn ctcaggctcc 300
ctancecett ecceptecet teccenecee egteceegee eegggggeeg eegecaeeeg 360
cctcccacca tggctctgaa ganaatccac aaggaattga a
<210> 264
<211> 401
<212> DNA
<213> Homo sapiens
<400> 264
aacaccagec actecaggac ceetgaagge etetaceagg teaceagtgt tetgegeeta 60
aagccacccc ctggcagaaa cttcagctgt gtgttctgga atactcacgt gagggaactt 120
actttggcca gcattgacct tcaaagtcag atggaaccca ggacccatcc aacttggctg 180
cttcacattt tcatcccctc ctgcatcatt gctttcattt tcatagccac agtgatagcc 240
ctaagaaaac aactetgtea aaagetgtat tetteaaaag acacaacaaa aagacetgte 300
accacaacaa agagggaagt gaacagtgct gtgaatctga acctgtggtc ttgggagcca 360
gggtgacctg atatgacatc taaagaagct tctggactct g
                                                                   401
<210> 265
<211> 271
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 59
<223> n = A, T, C or G
<400> 265
gccactteet gtggacatgg gcagageget gctgccagtt cctggtagec ttgaccacna 60
cgctgggggg tctttgtgat ggtcatgggt ctcatttgca cttgggggtg tgggattcaa 120
gttagaagtt totagatotg googggogoa gtggotoaca cotgtaatoo cagcacttta 180
ggaggctgag gcaggcggat catgaggtca ggagatcgag accgtcctgg ctaacacagt 240
gaaaccccgt ctctactaaa aatacaaaaa a
<210> 266
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 45
<223> n = A, T, C or G
<400> 266
attcataaat ttagctgaaa gatactgatt caatttgtat acagngaata taaatgagac 60
gacagcaaaa ttttcatgaa atgtaaaata tttttatagt ttgttcatac tatatgaggt 120
totattttaa atgactttot ggattttaaa aaatttottt aaatacaato atttttgtaa 180
tatttatttt atgcttatga tctagataat tgcagaatat cattttatct gactctgtct 240
tcataagaga gctgtggccg aattttgaac atctgttata gggagtgatc aaattagaag 300
gcaatgtgga aaaacaattc tgggaaagat ttctttatat gaagtccctg ccactagcca 360
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gccatcctaa ttgatgaaag ttatctgttc acaggcctgc a
                                                                   401
<210> 267
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 116, 247, 277, 296, 307, 313, 322, 323, 336, 342, 355, 365,
377, 378, 397
<223> n = A, T, C or G
<400> 267
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tgtggagtcg gatcctcttc ggggtgagcc agggtcggcg cgcgcggctg tctcanaact 120
catgcagctg ttcccgcgag gcctgtttga ggacgcgctg ccgcccatcg tgctgaggag 180
ccaggtqtac aqcettqtqc ctqacagqac cqtqqccqac cqqcaqctqa agqagcttca 240
agagcanggg gagacaaaat cgtccagctg ggcttcnact tggatgccca tggaanttat 300
tetttenett ganggaetta enngggaece aagaaneeet tneaagggge eettngtgga 360
tgggncccga aaccccnnta tttgcccttg ggggggncca a
                                                                   401
<210> 268
<211> 223
<212> DNA
<213> Homo sapiens
<400> 268
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ctcccaaagt gctgggatta caggtgtgag ccaccgcgcc tggcctgata catactttta 120
qaatcaaqta qtcacqcact ttttctqttc atttttctaa aaaqtaaata tacaaatqtt 180
ttgttttttg tttttttgt ttgtttgttt ctgttttttt ttt
                                                                   223
<210> 269
<211> 401
<212> DNA
<213> Homo sapiens
<400> 269
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tgctagttca tttgaatatt tctcccaact tatccaagga tctccagctc taacaaaatg 120
gtttattttt atttaaatgt caatagttgt tttttaaaat ccaaatcaga ggtgcaggcc 180
accagttaaa tgccgtctat caggttttgt gccttaagag actacagagt caaagctcat 240
ttttaaagga gtaggacaaa gttgtcacag gtttttgttg ttgtttttat tgcccccaaa 300
attacatgtt aatttecatt tatateaggg attetattta ettgaagaet gtgaagttge 360
cattttgtct cattgttttc tttgacataa ctaggatcca t
                                                                   401
<210> 270
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
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<222> 240, 382
<223> n = A,T,C or G
<400> 270
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ccttgtcaac tgaaaaatgc acctgacttc gagcaagact ctttccttag gttctggatc 120
tgtttgagcc ccatggcact gagctggaat ctgagggtct tgttccaagg atgtgatgat 180
gtgggagaat gttctttgaa agagcagaaa tccagtctgc atggaaacag cctgtagagn 240
agaagtttcc agtgataagt gttcactgtt ctaaggaggt acaccacagc tacctgaatt 300
tteccaaaat gagtgettet gtgegttaca actggeettt gtacttgact gtgatgaett 360
tgttttttct tttcaattct anatgaacat gggaaaaaat g
                                                                   401
<210> 271
<211> 329
<212> DNA
<213> Homo sapiens
<400> 271
ccacagcete caagteaggt ggggtggagt eccagagetg cacagggttt ggeecaagtt 60
tctaagggag gcacttcctc ccctcgccca tcagtgccag cccctgctgg ctggtgcctg 120
agreecteag acageceet geeegcagg cetgeettet cagggactte tgeggggeet 180
gaggcaagcc atggagtgag acccaggagc cggacacttc tcaggaaatg gcttttccca 240
acceccagee eccaeceggt ggttetteet gttetgtgae tgtgtatagt gecaecaeag 300
cttatggcat ctcattgagg acaaaaaa
                                                                   329
<210> 272
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 1, 7, 12, 21, 61, 62, 66, 72, 78, 88, 90, 92, 98, 117, 119,
128, 130, 134, 142, 144, 151, 159, 162, 164, 168, 169, 177,
184, 185, 188, 194, 202, 204, 209, 213, 218, 223, 231, 260,
272, 299, 300, 306, 321, 322, 323, 331, 335, 336, 338
<223> n = A, T, C or G
<221> misc feature
<222> 341, 342, 343, 345, 346, 351, 358, 360, 362, 363, 387, 390,
<223> n = A, T, C or G
<400> 272
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nncatnatat eneteatnge tgggeeentn angacaenat eccaetecaa eacetgngng 120
atgctggnen cetnggaace anenteagaa ngaccetgnt entntgtnnt eegeaanetg 180
aagnnaange gggntacace thentgeant ggnecaenet gengggaact ntacacacet 240
acgggatgtg gctgcgccan gagccaagag cntttctgga tgattcccca gcctcttgnn 300
aggganteta caacattget nnntacettt nteennenge nnntnntgga ntacaggngn 360
tnntaacact acatcttttt tactgeneen tnettggtgg g
                                                                   401
<210> 273
<211> 401
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<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 399
<223> n = A, T, C or G
<400> 273
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tggctccatc ctggcctcac tgtccacctt ccagcagatg tggattagca agcaggagta 120
cgacgagtcg ggcccctcca tcgtccaccg caaatgcttc taaacggact cagcagatgc 180
qtaqcatttq ctqcatqqqt taattqaqaa taqaaatttq cccctqqcaa atqcacacac 240
ctcatqctaq cctcacqaaa ctqqaataaq ccttcqaaaa gaaattqtcc ttqaaqcttq 300
tatctgatat cagcactgga ttgtagaact tgttgctgat tttgaccttg tattgaagtt 360
aactgttccc cttggtatta acgtgtcagg gctgagtgnt c
<210> 274
<211> 401
<212> DNA
<213> Homo sapiens
<400> 274
ccacccacac ccaccgcgcc ctcgttcgcc tcttctccgg gagccagtcc gcgccaccgc 60
egeogeocag gecategoca coeteegoag coatgteeae caggteegtg teetegteet 120
cctaccgcag gatgttcggc ggcccgggca ccgcgagccg gccgagctcc agccggagct 180
acgtgactac gtccaccegc acctacagec tgggcagegc getgegeece ageaccagec 240
geagesteta egestegtes segggeggeg tgtatgesas gegstestet geogtgeges 300
tgcggagcag cgtgcccggg gtgcggctcc tgcaggactc ggtggacttc tcgctggccg 360
acqccatcaa caccgagttc aagaacaccc gcaccaacga q
                                                                   401
<210> 275
<211> 401
<212> DNA
<213> Homo sapiens
<400> 275
ccacttccac cactttgtgg ageagtgeet teagegeaac eeggatgeea ggtateeetg 60
ctggcctggg cctgggcttc gggagagcag agggtgctca ggagggtaag gccagggtgt 120
gaagggactt acctcccaaa ggttctgcag gggaatctgg agctacacac aggagggatc 180
agctcctggg tgtgtcagag gccagcctgg ggagctctgg ccactgcttc ccatgagctg 240
agggagaggg agaggggacc cgaggctgag gcataagtgg caggatttcg ggaagctggg 300
gacacggcag tgatgctgcg gtctctcctc ccctttccct ccaggcccag tgccagcacc 360
ctcctgaacc actctttctt caagcagatc aagcgacgtg c
                                                                   401
<210> 276
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 11
<223> n = A, T, C or G
```

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<400> 276
totgatattg ntaccottga godacotaag ttagaagaaa ttggaaatca agaagttgto 60
attgttgaag aagcacagag ttcagaagac tttaacatgg gctcttcctc tagcagccag 120
tatactttct gtcagccaga aactgtattt tcatctcagc ctagtgatga tgaatcaagt 180
agtgatgaaa ccagtaatca geecagteet geetttagae gaegeegtge taggaagaag 240
accepttecte etteagaate teaagacege etagttegte aacaagaaac teaacettet 300
aaggagttga gtaaacgtca gttcagtagt ggtctcaata agtgtgttat acttgctttg 360
qtgattqcaa tcaqcatqqq atttqqccat ttctatqqca c
<210> 277
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 227, 333
<223> n = A, T, C or G
<400> 277
aactttggca acatatctca gcaaaaacta cagctatgtt attcatgcca aaataaaagc 60
tgtgcagagg agtggctgca atgaggtcac aacggtggtg gatgtaaaag agatcttcaa 120
gtoctcatca cocatocoto gaactcaagt cocgetcatt acaaattott ettgecagtg 180
tecacacate etgeeceate aagatgttet cateatgtgt tacgagngge geteaaggat 240
gatgettett gaaaattget tagttgaaaa atggagagat cagettagta aaagateeat 300
acagtgggaa gagaggctgc aggaacagcg ganaacagtt caggacaaga agaaaacagc 360
cgggcgcacc agtcgtagta atccccccaa accaaaggga a
                                                                   401
<210> 278
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 322, 354
<223> n = A, T, C or G
<400> 278
aatgagtgtg agaccacaaa tgaatgccgg gaggatgaaa tgtgttggaa ttatcatggc 60
ggetteegtt gttateeacg aaateettgt caagateeet acattetaac accagagaac 120
cgatgtgttt gcccagtctc aaatgccatg tgccgagaac tgccccagtc aatagtctac 180
aaatacatga gcatccgatc tgataggtct gtgccatcag acatcttcca gatacaggec 240
acaactattt atgccaacac catcaatact tttcggatta aatctggaaa tgaaaatgga 300
gagtetacet acgacaacaa anceetgtaa gtgcaatget tgtgetegtg aagneattat 360
caggaccaag agaacatatc gtggacctgg agatgctgac a
                                                                   401
<210> 279
<211> 401
<212> DNA
<213> Homo sapiens
<220>
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<221> misc feature
<222> 30, 35, 81, 88, 180, 212, 378, 384, 391
<223> n = A, T, C or G
<400> 279
aaattattgc ctctgataca tacctaagtn aacanaacat taatacctaa gtaaacataa 60
cattacttqq agggttqcaq nttctaantq aaactgtatt tgaaactttt aagtatactt 120
tagqaaacaa qcatqaacqq caqtctagaa taccagaaac atctacttgg gtagcttggn 180
qccattatcc tqtqqaatct gatatqtctg gnagcatgtc attgatggga catgaagaca 240
tetttggaaa tgatgagatt attteetgtg ttaaaaaaaa aaaaaatett aaatteetae 300
aatgtgaaac tgaaactaat aattttgatc ctgatgtatg ggacagcgta tctgtaccag 360
gctctaaata acaaaagnta gggngacaag nacatgttcc t
                                                                   401
<210> 280
<211> 326
<212> DNA
<213> Homo sapiens
<400> 280
gaagtggaat tgtataattc aattcgataa ttgatctcat gggctttccc tggaggaaag 60
gtttttttttt ttgtttttt tttaagaact tgaaacttgt aaactgagat gtctgtagct 120
tttttgccca tctgtagtgt atgtgaagat ttcaaaacct gagagcactt tttctttgtt 180
tagaattatg agaaaggcac tagatgactt taggatttgc atttttccct ttattgcctc 240
atttcttgtg acgccttgtt ggggagggaa atctgtttat tttttcctac aaataaaaag 300
                                                                   326
ctaagattct atatcgcaaa aaaaaa
<210> 281
<211> 374
<212> DNA
<213> Homo sapiens
<400> 281
caacgcqttt qcaaatattc ccctqqtaqc ctacttcctt acccccqaat attggtaaga 60
tegageaatg getteaggae atgggttete tteteetgtg ateatteaag tgeteaetge 120
atgaagactg gettgtetea gtgttteaac eteaceaggg etgtetettg gtecacacet 180
egeteeetgt tagtgeegta tgacageece catcaaatga eettggeeaa gteaeggttt 240
ctctqtqqtc aaqqttqqtt qqctqattqq tqqaaaqtaq qqtqqaccaa aggaggccac 300
gtgagcagtc agcaccagtt ctgcaccagc agcgcctccg tcctagtggg tgttcctgtt 360
teteetggee etgg
                                                                   374
<210> 282
<211> 404
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 26, 27, 51, 137, 180, 222
<223> n = A, T, C or G
<400> 282
agtgtggtgg aattcccgca tcctanncgc cgactcacac aaggcagagt ngccatggag 60
aaaattccag tgtcagcatt cttgctcctt gtggccctct cctacactct ggccagagat 120
accacaqtca aacctgnagc caaaaaggac acaaaggact ctcgacccaa actgccccan 180
```

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acceteteca gaggttgggg tgaccaacte atetggacte anacatatga agaageteta 240
tataaatcca agacaagcaa caaaccettg atgattatte atcaettgga tgagtgeeca 300
cacagtcaag ctttaaagaa agtgtttgct gaaaataaag aaatccagaa attggcagag 360
cagtttgtcc tcctcaatct ggtttatgaa acaactgaca aaca
<210> 283
<211> 184
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 26
<223> n = A, T, C or G
<400> 283
agtgtggtgg aattcacttg cttaanttgt gggcaaaaga gaaaaagaag gattgatcag 60
agcattgtgc aatacagttt cattaactcc ttccctcgct cccccaaaaa tttgaatttt 120
tttttcaaca ctcttacacc tgttatggaa aatgtcaacc tttgtaagaa aaccaaaata 180
<210> 284
<211> 421
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 147, 149
<223> n = A, T, C or G
<400> 284
ctattaatcc tgccacaata tttttaatta cgtacaaaga tctgacatgt cacccaggga 60
cccatttcac ccactgetet gtttggccgc cagtettttg tetetetet cagcaatggt 120
gaggeggata ecetteete ggggaanana aateeatggt ttgttgeeet tgeeaataae 180
aaaaatgttg gaaagtcgag tggcaaagct gttgccattg gcatctttca cgtgaaccac 240
gtcaaaagat ccagggtgcc tetetetgtt ggtgatcaca ccaattette etaggttage 300
acctccagtc accatacaca ggttaccagt gtcgaacttg atgaaatcag taatcttgcc 360
agtetetaaa teaatetgaa tggtateatt eacettgatg aggggategg ggtageggat 420
g
                                                                   421
<210> 285
<211> 361
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 34, 188
<223> n = A, T, C or G
<400> 285
ctgggtggta actctttatt tcattgtccg gaanaaagat gggagtggga acagggtgga 60
cactgtgcag gcttcagctt ccactccggg caggattcag gctatctggg accgcaggga 120
```

```
ctgccaggtg cacageeetg geteeegagg caggeaggea aggtgaeggg actggaagee 180
     cttttcanag ccttggagga gctggtccgt ccacaagcaa tgagtgccac tctgcagttt 240
     qcaqqqqatq qataaacaqq qaaacactgt gcatteetca cagecaacag tgtaggtett 300
     ggtgaagccc cggcgctgag ctaagctcag gctgttccag ggagccacga aactgcaggt 360
                                                                           361
     <210> 286
     <211> 336
     <212> DNA
     <213> Homo sapiens
     <220>
     <221> misc feature
     <222> 40, 68, 75, 127, 262
     <223> n = A, T, C or G
     <400> 286
     tttgagtggc agegeettta tttgtggggg cetteaaggn agggtegtgg ggggeagegg 60
     ggaggaanag ccganaaact gtgtgaccgg ggcctcaggt ggtgggcatt gggggctcct 120
11)
     cttgcanatg cccattggca tcaccggtgc agccattggt ggcagcgggt accggtcctt 180
     tcttgttcaa catagggtag gtggcagcca cgggtccaac tcgcttgagg ctgggccctg 240
ij
٠. ۽
پوءِ
     ggcgctccat tttgtgttcc angagcatgt ggttctgtgg cgggagcccc acgcaggccc 300
1,3
                                                                           336
     tgaggatgtt ctcgatgcag ctgcgctggc ggaaaa
٠...
     <210> 287
ij
     <211> 301
ï
     <212> DNA
135
     <213> Homo sapiens
<220>
III.
     <221> misc feature
     \langle 222 \rangle 15, \overline{3}3, 44, 53, 76, 83, 107, 117, 154, 166, 192, 194, 207,
E E
     215, 241, 246
ž = å:
     <223> n = A, T, C or G
     <400> 287
     tgggtaccaa atttntttat ttgaaggaat ggnacaaatc aaanaactta agnggatgtt 60
     ttggtacaac ttatanaaaa ggnaaaggaa accccaacat gcatgcnctg ccttggngac 120
     cagggaagtc accccacggc tatggggaaa ttancccgag gcttancttt cattatcact 180
     qtctcccaqq qnqnqcttqt caaaaanata ttccnccaag ccaaattcgg gcgctcccat 240
     nttgeneaag ttggteaegt ggteaeceaa ttetttgatg gettteaect geteatteag 300
                                                                           301
     <210> 288
     <211> 358
     <212> DNA
     <213> Homo sapiens
     <220>
     <221> misc feature
     <222> 39, 143, 226
     <223> n = A, T, C or G
     <400> 288
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aagtttttaa actttttatt tqcatattaa aaaaattqnq cattccaata attaaaatca 60
           tttgaacaaa aaaaaaaatg gcactctgat taaactgcat tacagcctgc aggacacctt 120
           gggccagctt ggttttactc tanatttcac tgtcgtccca ccccacttct tccaccccac 180
           ttcttccttc accaacatgc aagttctttc cttccctgcc agccanatag atagacagat 240
           gggaaaggca ggcgcgcct tcgttgtcag tagttctttg atgtgaaagg ggcagcacag 300
           tcatttaaac ttgatccaac ctctttgcat cttacaaagt taaacagcta aaagaagt
           <210> 289
           <211> 462
           <212> DNA
           <213> Homo sapiens
           <220>
           <221> misc feature
           <222> 87, 141, 182, 220, 269, 327
           <223> n = A, T, C or G
           <400> 289
and the state of t
           ggcatcagaa atgctgttta tttctctgct gctcccaagc tggctggcct ttgcagagga 60
111
           gcagacaaca gatgcatagt tgggganaaa gggaggacag gttccaggat agagggtgca 120
ď
           ggctgaggga ggaagggtaa naggaaggaa ggccatcctg gatccccaca tttcagtctc 180
           anatgaggac aaagggactc ccaagccccc aaatcatcan aaaacaccaa ggagcaggag 240
1,5
           gagettgage aggeeccagg gageeteana geeataccag ecaetgteta etteccatee 300
tectetecca ttecetgtet getteanace accteecage taageceeag etecatteee 360
1,2
           ccaatectgg ceettgecag ettgacagte acagtgectg gaattecace actgaggett 420
100
           ctcccagttg gattaggacg tcgccctgtt agcatgctgc cc
                                                                                                                                                       462
ij
<210> 290
122
           <211> 481
           <212> DNA
<213> Homo sapiens
112
<220>
i ak
           <221> misc feature
           <222> 44, 57, 122, 158, 304, 325, 352, 405
           <223> n = A, T, C or G
           <400> 290
           tactttccta aactttatta aagaaaaaag caataagcaa tggnggtaaa tctctanaac 60
           atacccaatt ttctgggctt cctcccccga gaatgtgaca ttttgatttc caaacatgcc 120
           anaagtgtat ggttcccaac tgtactaaag taggtganaa gctgaagtcc tcaagtgttc 180
           atcttccaac ttttcccagt ctgtggtctg tctttggatc agcaataatt gcctgaacag 240
           ctactatggc ttcgttgatt tttgtctgta gctctctgag ctcctctatg tgcagcaatc 300
           gcanaatttg agcagettea ttaanaactg cateteetgt gteaaaaeca anaatatgtt 360
           tgtctaaagc aacaggtaag coctcttttg tttgatttgc cttancaact gcatcctgtg 420
           teaggegete etgaaceaaa ateegaattg cettaageat taeeaggtaa teateatgae 480
                                                                                                                                                       481
           <210> 291
           <211> 381
           <212> DNA
           <213> Homo sapiens
           <220>
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<221> misc feature
<222> 79, 166, 187, 208, 219, 315
<223> n = A, T, C or G
<400> 291
tcatagtaat gtaaaaccat ttgtttaatt ctaaatcaaa tcactttcac aacagtgaaa 60
attagtgact ggttaaggng tgccactgta catatcatca ttttctgact ggggtcagga 120
cctqqtccta qtccacaaqq qtqqcaqqaq gagggtggag gctaanaaca cagaaaacac 180
acaaaanaaa qqaaaqctqc cttqqcanaa ggatgaggng gtgagcttgc cgaaggatgg 240
tgggaagggg geteeetgtt ggggeegage eaggagteee aagteagete teetgeetta 300
cttagctcct ggcanagggt gagtggggac ctacgaggtt caaaatcaaa tggcatttgg 360
                                                                   381
ccagcctggc tttactaaca g
<210> 292
<211> 371
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 32, 55, 72, 151, 189, 292
<223> n = A, T, C or G
<400> 292
qaaaaaataa teeqtttaat tgaaaaacet gnaggataet attecaetee eecanatgag 60
gaggetgagg anaccaaacc cetacateac etegtageca ettetgatac tetteacgag 120
gcagcaggca aagacaattc ccaaaacctc nacaaaagca attccaaggg ctgctgcagc 180
taccaccanc acatttttcc tcagccagcc cccaatcttc tccacacagc cctccttatg 240
qatcqccttc tcgttgaaat taatcccaca gcccacagta acattaatgc ancaggagtc 300
ggqqactcqq ttcttcqaca tggaagggat tttctcccaa tctgtgtagt tagcagcccc 360
                                                                   371
acaqcactta a
<210> 293
<211> 361
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 75, 196, 222
<223> n = A, T, C or G
<400> 293
qatttaaaaq aaaacacttt attgttcagc aattaaaagt tagccaaata tgtatttttc 60
tccataattt attgngatgt tatcaacatc aagtaaaatg ctcattttca tcatttgctt 120
ctqttcatqt tttcttgaac acqtcttcaa ttttccttcc aaaatgctgc atgccacact 180
tgaggtaacg aagcanaagt atttttaaac atgacagcta anaacattca tctacagcaa 240
cctatatgct caatacatgc cgcgtgatcc tagtagtttt ttcacaacct tctacaagtt 300
tttggaaaac atctgttatg atgactttca tacaccttca cctcaaaggc tttcttgcac 360
                                                                    361
<210> 294
<211> 391
<212> DNA
```

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<212> DNA

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<213> Homo sapiens
<220>
<221> misc feature
\langle 222 \rangle 26, \overline{7}7, 96, 150, 203, 252, 254, 264, 276
<223> n = A, T, C or G
<400> 294
tattttaaaq tttaattatq attcanaaaa aatcqaqcqa ataactttct ctgaaaaaat 60
atattgactc tgtatanacc acagttattg gggganaagg gctggtaggt taaattatcc 120
tattttttat tctgaaaatg atattaatan aaagtcccgt ttccagtctg attataaaga 180
tacatatqcc caaaatqqct qanaataaat acaacaggaa atqcaaaagc tqtaaaqcta 240
agggcatgca ananaaaatc tcanaatacc caaagnggca acaaggaacg tttggctgga 300
atttgaagtt atttcagtca tetttgtett tggeteeatg tttcaggatg egtgtgaact 360
cgatgtaatt gaaattcccc tttttatcaa t
<210> 295
<211> 343
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 145, 174, 205, 232
<223> n = A, T, C or G
<400> 295
ttettttgtt ttattgataa cagaaactgt geataattae agatttgatg aggaatetge 60
aaataataaa gaatgtgtct actgccagca aaatacaatt attccatgcc ctctcaacat 120
acaaatataq aqttcttcac accanatggc tctggtgtaa caaagccatt ttanatgttt 180
aattgtgctt ctacaaaacc ttcanagcat gaggtagttt cttttaccta cnatattttc 240
cacatttcca ttattacact tttagtgagc taaaatcctt ttaacatagc ctgcggatga 300
tctttcacaa aagccaagcc tcatttacaa agggtttatt tct
                                                                     343
<210> 296
<211> 241
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 96, 98, 106, 185
<223> n = A, T, C or G
<400> 296
ttcttggata ttggttgttt ttgtgaaaaa gtttttgttt ttcttctcag tcaactgaat 60
tatttctcta ctttgccctc ctgatgccca catgananaa cttaanataa tttctaacag 120
cttccacttt ggaaaaaaa aaaacctgtt ttcctcatgg aaccccagga gttgaaagtg 180
gatanatogo totoaaaato taaggototg ttoagottta cattatgtta cotgacgttt 240
                                                                     241
<210> 297
<211> 391
```

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the first care care care care as a care of a fact of the care of t
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<213> Homo sapiens
<220>
<221> misc feature
<222> 12, 130
\langle 223 \rangle n = A,T,C or G
<400> 297
gttgtggctg anaatgctgg agatgctcag ttctctccct cacaaggtag gccacaaatt 60
cttggtggtg ccctcacatc tggggtcttc aggcaccagc catgcctgcc gaggagtgct 120
gtcaggacan accatgtccg tgctaggccc aggcacagcc caaccactcc tcatccaagt 180
ctctcccagg tttctggtcc cgatgggcaa ggatgacccc tccagtggct ggtaccccac 240
cateceacta ecceteacat geteteacte tecateaggt ecceaateet ggetteecte 300
ttcacqaact ctcaaagaaa aggaaggata aaacctaaat aaaccagaca gaagcagctc 360
tggaaaagta caaaaagaca gccagaggtg t
<210> 298
<211> 321
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 14, 30, 76, 116, 201, 288, 301
<223> n = A, T, C or G
<400> 298
caagccaaac tgtntccagc tttattaaan atactttcca taaacaatca tggtatttca 60
ggcaggacat gggcanacaa tcgttaacag tatacaacaa ctttcaaact cccttnttca 120
atggactacc aaaaatcaaa aagccactat aaaacccaat gaagtettea tetgatgete 180
tgaacaggga aagtttaaag ngagggttga catttcacat ttagcatgtt gtttaacaac 240
ttttcacaag ccqaccctqa ctttcaqqaa qtqaaatqaa aatgqcanaa tttatctqaa 300
natccacaat ctaaaaatgg a
                                                                    321
<210> 299
<211> 401
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 104, 268, 347
<223> n = A, T, C or G
<400> 299
tatcataaag agtgttgaag tttatttatt atagcaccat tgagacattt tgaaattgga 60
attggtaaaa aaataaaaca aaaagcattt gaattgtatt tggnggaaca gcaaaaaaag 120
agaagtatca tttttctttg tcaaattata ctgtttccaa acattttgga aataaataac 180
tggaattttg tcggtcactt gcactggttg acaagattag aacaagagga acacatatgg 240
agttaaattt tttttgttgg gatttcanat agagtttggt ttataaaaag caaacagggc 300
caacqtccac accaaattct tqatcagqac caccaatqtc ataqqqnqca atatctacaa 360
taggtagtct cacagccttq cqtqttcqat attcaaaqac t
                                                                    401
```

<210> 300

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SHEAR CHIEF, ALTHOUGH STEIN, SHEEN, THE SHEEP, ST. TO THE THE SHEEP, SHE
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<211> 188
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 48
<223> n = A, T, C or G
<400> 300
tqaatqcttt qtcatattaa qaaaqttaaa qtqcaataat qtttqaanac aataagtggt 60
ggtgtatctt gtttctaata agataaactt ttttgtcttt gctttatctt attagggagt 120
tgtatgtcag tgtataaaac atactgtgtg gtataacagg cttaataaat tctttaaaag 180
                                                                   188
qaaaaaaa
<210> 301
<211> 291
<212> DNA
<213> Homo sapiens
<400> 301
aagattttgt tttattttat tatggctaga aagacactgt tatagccaaa atcggcaatg 60
acactaaaga aatcctctgt gcttttcaat atgcaaatat atttcttcca agagttgccc 120
tggtgtgact tcaagagttc atgttaactt cttttctgga aacttccttt tcttagttgt 180
tgtattettg aagageetgg gecatgaaga gettgeetaa gttttgggea gtgaacteet 240
                                                                   291
tgatgttctg gcagtaagtg tttatctggc ctgcaatgag cagcgagtcc a
<210> 302
<211> 341
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 25
<223> n = A, T, C or G
<400> 302
tgatttttca taattttatt aaatnatcac tgggaaaact aatggttcgc gtatcacaca 60
attacactac aatctgatag gagtggtaaa accagccaat ggaatccagg taaagtacaa 120
aaacgccacc ttttattgtc ctgtcttatt tctcgggaag gagggttcta ctttacacat 180
ttcatgagcc agcagtggac ttgagttaca atgtgtaggt tccttgtggt tatagctgca 240
gaaqaaqcca tcaaattctt qaqqacttqa catctctcqq aaaqaaqcaa actaqtqqat 300
ccccgggct gcaggaattc gatatcaagc ttatcgatac c
                                                                    341
<210> 303
<211> 361
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 15, 27, 92, 124, 127, 183, 198, 244, 320
<223> n = A, T, C or G
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<400> 303
tgcagacagt aaatnaattt tatttgngtt cacagaacat actaggcgat ctcgacagtc 60
getecqtqae ageceaceaa eccecaacee thtacetege agecaceeta aaggegaett 120
caanaanatg gaaggatete aeggatetea tteetaatgg teegeegaag teteacaeag 180
tanacagacg gagttganat gctggaggat gcagtcacct cctaaactta cgacccacca 240
ccanacttca teccageegg gaegteetee eccaeeegag teeteeceat ttetteteet 300
actttgccgc agttccaggn gtcctgcttc caccagtccc acaaagctca ataaatacca 360
<210> 304
<211> 301
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 23, 104, 192
<223> n = A, T, C or G
<400> 304
ctctttacaa cagcctttat ttncggccct tgatcctgct cggatgctgg tggaggccct 60
tageteegee egecaggete tgtgeegeet eeeegeagge geanatteat gaacaeggtg 120
ctcaggggct tgaggccgta ctccccagc gggagctggt cctccagggg cttcccctcg 180
aaggtcagcc anaacaggtc gtcctgcaca ccctccagcc cgctcacttg ctgcttcagg 240
tgggccacgg tctgcgtcag ccgcacctcg taggtgctgc tgcggccctt gttattcctc 300
                                                                   301
<210> 305
<211> 331
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 3, 36, 60, 193, 223
<223> n = A, T, C or G
<400> 305
qanaqqctaq taacatcagt tttattgggt tggggnggca accatagcct ggctgggggn 60
ggggctggcc ctcacaggtt gttgagttcc agcagggtct ggtccaaggt ctggtgaatc 120
tegaegttet eeteettgge aetggeeaag gtetetteta ggteategat ggttttetee 180
aactttgcca canacctctc ggcaaactct gctcgggtct cancetcctt cagcttctcc 240
tccaacagtt tgatctcctc ttcatattta tcttctttgg gggaatactc ctcctctgag 300
                                                                   331
gccatcaggg acttgagggc ctggtccatg g
<210> 306
<211> 457
<212> DNA
<213> Homo sapiens
<400> 306
aatatgtaaa ggtaataact tttattatat taaagacaat gcaaacgaaa aacagaattg 60
agcagtgcaa aatttaaagg actgttttgt tctcaaagtt gcaagtttca aagccaaaag 120
```

```
aattatatgt atcaaatata taagtaaaaa aaagttagac tttcaagcct gtaatcccag 180
cactttggga ggctgaggca ggtggatcac taacattaaa aagacaacat tagattttgt 240
cgatttatag caattttata aatatataac tttgtcactt ggatcctgaa gcaaaataat 300
aaagtgaatt tgggattttt gtacttggta aaaagtttaa caccctaaat tcacaactag 360
tggatccccc gggctgcagg aattcgatat caagcttatc gataccgtcg acctcgaggg 420
ggggcccggt acccaattcg ccctatagtg agtcgta
                                                                   457
<210> 307
<211> 491
<212> DNA
<213> Homo sapiens
<400> 307
gtgettggae ggaaceegge getegtteee caceeeggee ggeegeecat ageeageect 60
ccgtcacctc ttcaccgcac cctcggactg ccccaaggcc cccgccgccg ctccagcgcc 120
gegeageeae egeegeegee geegeetete ettagtegee geeatgaega eegegteeae 180
ctcgcaggtg cgccagaact accaccagga ctcagaggcc gccatcaacc gccagatcaa 240
cctggagete tacgeetect acgtttacet gtecatgtet tactactttg accgegatga 300
tgtggctttg aagaactttg ccaaatactt tcttcaccaa tctcatgagg agagggaaca 360
tgctgagaaa ctgatgaagc tgcagaacca acgaggtggc cgaatcttcc ttcaggatat 420
caagaaacca gactgtgatg actgggagag cgggctgaat gcaatggagt gtgcattaca 480
tttggaaaaa a
                                                                   491
<210> 308
<211> 421
<212> DNA
<213> Homo sapiens
<400> 308
ctcagcgctt cttctttctt ggtttgatcc tgactgctgt catggcgtgc cctctggaga 60
aggccctgga tgtgatggtg tccaccttcc acaagtactc gggcaaagag ggtgacaagt 120
tcaagctcaa caagtcagaa ctaaaggagc tgctgacccg ggagctgccc agcttcttgg 180
ggaaaaggac agatgaagct gctttccaga agctgatgag caacttggac agcaacaggg 240
acaacgaggt ggacttccaa gagtactgtg tetteetgte etgeategee atgatgtgta 300
acgaattett tgaaggette eeagataage ageecaggaa gaaatgaaaa eteetetgat 360
gtggttgggg ggtctgccag ctggggccct ccctgtcgcc agtgggcact tttttttttc 420
<210> 309
<211> 321
<212> DNA
<213> Homo sapiens
<400> 309
accaaatggc ggatgacgcc ggtgcagcgg gggggcccgg gggccctggt ggccctggga 60
tggggaaceg eggtggette egeggaggtt teggeagtgg cateegggge eggggtegeg 120
geegtggaeg gggeegggge egaggeegeg gagetegegg aggeaaggee gaggataagg 180
agtggatgcc cgtcaccaag ttgggccgct tggtcaagga catgaagatc aagtccctgg 240
aggagateta tetettetee etgeceatta aggaateaga gateattgat ttetteetgg 300
gggcctctct caaggatgag g
                                                                   321
<210> 310
<211> 381
<212> DNA
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<213> Homo sapiens
<400> 310
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tcagtgccta tttttcctgg aaactcaatt ttaaatagtc caattccatc tgaagccaag 120
ctgttgtcat tttcattcgg tgacattctc tcccatgaca cccagaaggg gcagaagaac 180
cacatttttc atttatagat gtttgcatcc tttgtattaa aattattttg aaggggttgc 240
ctcattggat ggcttttttt tttttcctcc agggagaagg ggagaaatgt acttggaaat 300
taatgtatgt ttacatctct ttgcaaattc ctgtacatag agatatattt tttaagtgtg 360
                                                                   381
aatgtaacaa catactgtga a
<210> 311
<211> 538
<212> DNA
<213> Homo sapiens
<400> 311
tttgaattta caccaagaac ttctcaataa aagaaaatca tgaatgctcc acaatttcaa 60
cataccacaa gagaagttaa tttcttaaca ttgtgttcta tgattatttg taagaccttc 120
accaagttct gatatctttt aaagacatag ttcaaaattg cttttgaaaa tctgtattct 180
tgaaaatatc cttgttgtgt attaggtttt taaataccag ctaaaggatt acctcactga 240
qtcatcaqta ccctcctatt cagetcccca agatgatgtg tttttgctta ccctaagaga 300
ggttttcttc ttatttttag ataattcaag tgcttagata aattatgttt tctttaagtg 360
tttatggtaa actcttttaa agaaaattta atatgttata getgaatett tttggtaaet 420
ttaaatettt ateatagaet etgtaeatat gtteaaatta getgettgee tgatgtgtgt 480
atcatcggtg ggatgacaga acaaacatat ttatgatcat gaataatgtg ctttgtaa
<210> 312
<211> 176
<212> DNA
<213> Homo sapiens
<400> 312
qqaqqaqcaq ctgaqaqata gggtcagtga atgcggttca gcctgctacc tctcctgtct 60
tcatagaacc attgccttag aattattgta tgacacgttt tttgttggtt aagctgtaag 120
qttttqttct ttqtqaacat gggtattttq aggggagggt ggagggagta gggaag
<210> 313
<211> 396
<212> DNA
<213> Homo sapiens
<400> 313
ccagcacccc caggccctgg gggacctggg ttctcagact gccaaagaag ccttgccatc 60
tggcgctccc atggctcttg caacatctcc ccttcgtttt tgagggggtc atgccggggg 120
agccaccage ceeteactgg gtteggagga gagteaggaa gggeeaagea egaeaaagea 180
gaaacatcgg atttggggaa cgcgtgtcaa tcccttgtgc cgcagggctg ggcgggagag 240
actgttctgt tccttgtgta actgtgttgc tgaaagacta cctcgttctt gtcttgatgt 300
gtcaccgggg caactgcctg ggggcgggga tggggggagg gtggaagcgg ctccccattt 360
                                                                   396
tataccaaag gtgctacatc tatgtgatgg gtgggg
<210> 314
<211> 311
<212> DNA
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<213> Homo sapiens
<400> 314
ceteaacate etcagagagg actggaagee agteettaeg ataaacteea taatttatgg 60
cctgcagtat ctcttcttgg agcccaaccc cgaggaccca ctgaacaagg aggccgcaga 120
ggtcctgcag aacaaccggc ggctgtttga gcagaacgtg cagcgctcca tgcggggtgg 180
ctacatcqqc tocacctact ttqaqcqctq cctqaaataq ggttggcgca tacccacccc 240
cgccacggcc acaagccctg gcatcccctg caaatattta ttgggggcca tgggtagggg 300
tttggggggc g
<210> 315
<211> 336
<212> DNA
<213> Homo sapiens
<400> 315
tttagaacat qqttatcatc caagactact ctaccctgca acattgaact cccaagagca 60
aatccacatt cctcttgagt tctgcagctt ctgtgtaaat agggcagctg tcgtctatgc 120
cqtaqaatca catqatctqa qqaccattca tqqaaqctqc taaataqcct aqtctgqgga 180
gtcttccata aagttttgca tggagcaaac aaacaggatt aaactaggtt tggttccttc 240
agecetetaa aageataggg ettageetge aggetteett gggetttete tgtgtgtgta 300
qttttqtaaa cactataqca tctgttaaga tccagt
<210> 316
<211> 436
<212> DNA
<213> Homo sapiens
<400> 316
aacatggtct gcgtgcctta agagagacgc ttcctgcaga acaggacctg actacaaaga 60
atgtttccat tqqaattqtt qqtaaaqact tqqaqtttac aatctatqat gatgatgatg 120
tgtctccatt cctgqaaqqt cttqaaqaaa gaccacagag aaaggcacag cctgctcaac 180
ctgctgatga acctgcagaa aaggctgatg aaccaatgga acattaagtg ataagccagt 240
ctatatatqt attatcaaat atqtaaqaat acaggcacca catactgatg acaataatct 300
atactttgaa ccaaaagttg cagagtggtg gaatgctatg ttttaggaat cagtccagat 360
gtgagttttt tecaagcaac eteaetgaaa eetatataat ggaatacatt tttetttgaa 420
                                                                   436
agggtctgta taatca
<210> 317
<211> 196
<212> DNA
<213> Homo sapiens
<400> 317
tattccttgt gaagatgata tactattttt gttaagcgtg tctgtattta tgtgtgagga 60
gctgctggct tgcagtgcgc gtgcacgtgg agagctggtg cccggagatt ggacggcctg 120
atgetecete ecetgeeetg gteeagggaa getggeegag ggteetgget eetgagggge 180
                                                                   196
atctgcccct ccccca
<210> 318
<211> 381
<212> DNA
<213> Homo sapiens
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<220>
<221> misc feature
<222> 8, 9, 102, 122, 167, 182, 193, 235, 253, 265, 266, 290, 321,
<223> n = A, T, C or G
<400> 318
gacgettnng cegtaacgat gateggagae ateetgetgt tegggaegtt getgatgaat 60
gccggggcgg tgctgaactt taagctgaaa aagaaggaca cncagggctt tggggaggag 120
thragggage ccaacacagg tgacaacate egggaattet tgetganeet cagataettt 180
cnaatettea tenecetgtg gaacatette atgatgttet geatgattgt getgntegge 240
tettgaatee canegatgaa accannaact caettteeeg ggatgeegan tetecattee 300
tocattoctg atgacttoaa naatgttttt gaccaaaaaa cogacaacct toccagaaag 360
                                                                   381
tccaaqctcq tqqtqqqnqq a
<210> 319
<211> 506
<212> DNA
<213> Homo sapiens
<400> 319
ctaagcttta cgaatggggt gacaacttat gataaaaact agagctagtg aattagccta 60
tttgtaaata cctttgttat aattgatagg atacatcttg gacatggaat tgttaagcca 120
cctctqaqca qtqtatqtca qqacttqttc attaggttgg cagcagaggg gcagaaggaa 180
ttatacaggt agagatgtat gcagatgtgt ccatatatgt ccatatttac attttgatag 240
ccattgatgt atgcatctct tggctgtact ataagaacac attaattcaa tggaaataca 300
ctttgctaat attttaatgg tatagatctg ctaatgaatt ctcttaaaaa catactgtat 360
totgttgctg tgtgtttcat tttaaattga gcattaaggg aatgcagcat ttaaatcaga 420
actotyccaa tyettttate tagaggegtg ttgccatttt tytettatat gaaatttetg 480
                                                                   506
teccaagaaa ggeaggatta catett
<210> 320
<211> 351
<212> DNA
<213> Homo sapiens
<400> 320
ctgacctgca ggacgaaacc atgaagagcc tgatccttct tgccatcctg gccgccttag 60
cggtagtaac tttgtgttat gaatcacatg aaagcatgga atcttatgaa cttaatccct 120
tcattaacag gagaaatgca aatacettca tateeeetca geagagatgg agagetaaag 180
tocaagagag gatoogagaa ogototaago otgtocacga gotoaatagg gaagootgtg 240
atgactacag actttgcgaa cgctacgcca tggtttatgg atacaatgct gcctataatc 300
                                                                   351
gctacttcag gaagcgccga gggaccaaat gagactgagg gaagaaaaaa a
<210> 321
<211> 421
<212> DNA
<213> Homo sapiens
<400> 321
ctcqqaqqcq ttcaqctqct tcaaqatqaa gctgaacatc tccttcccag ccactggctg 60
ccagaaactc attgaagtgg acgatgaacg caaacttcgt actttctatg agaagcgtat 120
ggccacagaa gttgctgctg acgctctggg tgaagaatgg aagggttatg tggtccgaat 180
caqtqqtqqq aacgacaaac aaggtttccc catgaagcag ggtgtcttga cccatggccg 240
```

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tgtccgcctg ctactgagta aggggcattc ctgttacaga ccaaggagaa ctggagaaag 300
aaagagaaaa teagttegtg gttgeattgt ggatgeaaat etgagegtte teaacttggt 360
tattqtaaaa aaaggagaga aggatattee tggaetgaet gataetaeag tgeetegeeg 420
                                                                   421
<210> 322
<211> 521
<212> DNA
<213> Homo sapiens
<400> 322
ageagetete etgecacage tecteacece etgaaaatgt tegeetgete caagtttgte 60
tecaetecet cettggteaa gageacetea eagetgetga geegteeget atetgeagtg 120
qtqctqaaac gaccgqaqat actgacaqat qaqaqcctca gcaqcttggc agtctcatgt 180
coccttacct cacttgtctc tagccgcage ttccaaacca gegecattte aagggacate 240
gacacagcag ccaagttcat tggagctggg gctgccacag ttggggtggc tggttctggg 300
gctgggattg gaactgtgtt tgggagcctc atcattggtt atgccaggaa cccttctctg 360
aagcaacage tetteteeta egecattetg ggetttgeee teteggagge catggggete 420
ttttgtctga tggtagcctt tctcatcctc tttgccatgt gaaggagccg tctccacctc 480
ccatagttct eccgcgtctg gttggccccg tgtgttcctt t
<210> 323
<211> 435
<212> DNA
<213> Homo sapiens
<400> 323
cogagging acquirigada etterecque deagacqued cogedatque etacquede 60
tectacetge tggetgeeet agggggeaac tecteeecca gegeeaagga catcaagaag 120
atottggaca gegtgggtat egaggeggae gaegaeegge teaacaaggt tateagtgag 180
ctgaatggaa aaaacattga agacgtcatt gcccagggta ttggcaagct tgccagtgta 240
cetgetggtg gggetgtage egtetetget geceeagget etgeageece tgetgetggt 300
tctgcccctg ctgcagcaga ggagaagaaa gatgagaaga aggaggagtc tgaagagtca 360
gatgatgaca tgggatttgg cctttttgat taaattcctg ctcccctgca aataaagcct 420
ttttacacat ctcaa
                                                                   435
<210> 324
<211> 521
<212> DNA
<213> Homo sapiens
<400> 324
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tggtgcagta caagaatcgt caggccatcc tggcggtcaa atccacgcgg cagaagcagc 120
agcacctggt ccagcagcag ccccctcgc agccgcagcc gcagccgcag ctccagcccc 180
aaccccagec teagecteag eegeaaceee ageceeaate acaaccccag ceteageeee 240
aacccaagec teageeceag eageteeace egtateegea tecacateea eatecacaet 300
cteatectea etegeaceea cacceteace egeaceegea teegeaceaa atacegeace 360
cacacccaca geogeacteg cageegeacg ggeacegget teteegeage acetecaaet 420
ctgcctgaaa ggggcagctc ccgggcaaga caaggttttg aggacttgag gaagtgggac 480
                                                                   521
gagcacattt ctattqtctt cacttggatc aaaagcaaaa c
<210> 325
<211> 451
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```
<212> DNA
<213> Homo sapiens
<400> 325
attttcattt ccattaacct ggaagctttc atgaatattc tcttctttta aaacatttta 60
acattattta aacagaaaaa gatgggctct ttctggttag ttgttacatg atagcagaga 120
tatttttact tagattactt tgggaatgag agattgttgt cttgaactct ggcactgtac 180
agtgaatgtg totgtagttg tgttagtttg cattaagcat gtataacatt caagtatgto 240
atccaaataa gaggcatata cattgaattg tttttaatcc tctgacaagt tgactcttcg 300
acccccaccc ccacccaaga cattttaata gtaaatagag agagagagaa gagttaatga 360
acatgaggta gtgttccact ggcaggatga cttttcaata gctcaaatca atttcagtgc 420
ctttatcact tgaattatta acttaatttg a
                                                                   451
<210> 326
<211> 421
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 296
<223> n = A, T, C or G
<400> 326
cgcggtcgta agggctgagg atttttggtc cgcacgctcc tgctcctgac tcaccgctgt 60
tegetetege egaggaacaa gteggteagg aageeegege geaacageea tggettttaa 120
ggatacegga aaaacaceeg tggageegga ggtggeaatt cacegaatte gaateaceet 180
aacaagccgc aacgtaaaat ccttggaaaa ggtgtgtgct gacttgataa gaggcgcaaa 240
agaaaagaat ctcaaagtga aaggaccagt tcgaatgcct accaagactt tgagantcac 300
tacaagaaaa actccttqtq qtqaaqqttc taagacqtqq qatcqtttcc aqatqagaat 360
tcacaagcga ctcattgact tgcacagtcc ttctgagatt gttaagcaga ttacttccat 420
                                                                   421
<210> 327
<211> 456
<212> DNA
<213> Homo sapiens
<400> 327
atettgaega ggetgeggtg tetgetgeta tteteegage ttegeaatge egeetaagga 60
cgacaagaag aagaaggacg ctggaaagtc ggccaagaaa gacaaagacc cagtgaacaa 120
atcogggggc aaggccaaaa agaagaagtg gtccaaaggc aaagttcggg acaagctcaa 180
taacttagtc ttgtttgaca aagctaccta tgataaactc tgtaaggaag ttcccaacta 240
taaacttata accccagctg tggtctctga gagactgaag attcgaggct ccctggccag 300
ggcagccctt caggagctcc ttagtaaagg acttatcaaa ctggtttcaa agcacagagc 360
tcaagtaatt tacaccagaa ataccaaggg tggagatgct ccagctgctg gtgaagatgc 420
atgaataggt ccaaccagct gtacatttgg aaaaat
                                                                   456
<210> 328
<211> 471
<212> DNA
<213> Homo sapiens
<400> 328
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gtggaagtga catcgtcttt aaaccctgcg tggcaatccc tgacgcaccg ccgtgatgcc 60
cagggaagac agggcgacct ggaagtccaa ctacttcctt aagatcatcc aactattgga 120
tgattatoog aaatgtttoa ttgtgggago agacaatgtg ggotocaago agatgcagca 180
gateegeatg teeettegeg ggaaggetgt ggtgetgatg ggeaagaaca ceatgatgeg 240
caaggccatc cgagggcacc tggaaaacaa cccagctctg gagaaactgc tgcctcatat 300
ccgggggaat gtgggctttg tgttcaccaa ggaggacctc actgagatca gggacatgtt 360
gctggccaat aaggtgccag ctgctgcccg tgctggtgcc attgccccat gtgaagtcac 420
tgtgccagcc cagaacactg gtctcgggcc cgagaagacc tcctttttcc a
<210> 329
<211> 278
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> 154, 204
<223> n = A, T, C \text{ or } G
<400> 329
gtttaaactt aagettggta cegagetegg atceactagt ceagtgtggt ggaattetag 60
aaattgagat gccccccag gccagcaaat gttccttttt gttcaaagtc tatttttatt 120
ccttgatatt tttcttttt ttttttttt ttgnggatgg ggacttgtga atttttctaa 180
aggtgctatt taacatggga gganagcgtg tgcggctcca gcccagcccg ctgctcactt 240
tecaecetet etecaectge etetggette teaggeet
                                                                 278
<210> 330
<211> 338
<212> DNA
<213> Homo sapiens
<400> 330
ctcaggette aacategaat aegeegeagg eccettegee etattettea tageegaata 60
cacaaacatt attataataa acacceteae cactacaate tteetaggaa caacatatga 120
egeactetee cetgaactet acacaacata ttttgtcace aagaceetae ttctaacete 180
cctgttctta tgaattcgaa cagcataccc ccgattccqc tacqaccaac tcatacacct 240
cattacaatc tccagcattc cccctcaaac ctaaaaaa
                                                                 338
<210> 331
<211> 2820
<212> DNA
<213> Homo sapiens
<400> 331
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ccatcagaag atggtgcgac aaacaagatt gagattagca tggactgtat ccqcatqcag 180
gactoggaco tgagtgacco catgtggcca cagtacacga acctggggct cotgaacago 240
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cagaacageg teaeggegee etegecetae geacageeea geteeacett egatgetete 360
tetecateae eegecateee etecaaeaee gaetaeeeag geeegeaeag titegaegig 420
tecttecage agtegageae egecaagteg gecaectgga egtattecae tgaactgaag 480
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Val Thr Ala Pro Ser Pro Tyr Ala Gln Pro Ser Pro Thr Phe Asp Ala
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Lys Thr Cys Pro Ile Gln Ile Lys Val Met Thr Pro Pro Pro Gln Gly
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Glu Val Val Lys Arg Cys Pro Asn His Glu Leu Ser Arg Glu Phe Asn
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Glu Gly Gln Ile Ala Pro Pro Ser His Leu Ile Arg Val Glu Gly Asn
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Ser His Ala Gln Tyr Val Glu Asp Pro Ile Thr Gly Arg Gln Ser Val
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Leu Val Pro Tyr Glu Pro Pro Gln Val Gly Thr Glu Phe Thr Thr Val
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                                                 205
Leu Tyr Asn Phe Met Cys Asn Ser Ser Cys Val Gly Gly Met Asn Arg
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Leu Gly Arg Arg Cys Phe Glu Ala Arg Ile Cys Ala Cys Pro Gly Arg
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Asp Arg Lys Ala Asp Glu Asp Ser Ile Arg Lys Gln Gln Val Ser Asp
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Ser Thr Lys Asn Gly Asp Gly Thr Lys Arg Pro Phe Arg Gln Asn Thr
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His Gly Ile Gln Met Thr Ser Ile Lys Lys Arg Arg Ser Pro Asp Asp
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Glu Leu Leu Tyr Leu Pro Val Arg Gly Arg Glu Thr Tyr Glu Met Leu
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Thr Ile Glu Thr Tyr Arg Gln Gln Gln Gln Gln His Gln His Leu
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Ile Pro Asp Gly Met Gly Ala Asn Ile Pro Met Met Gly Thr His Met
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Pro Met Ala Gly Asp Met Asn Gly Leu Ser Pro Thr Gln Ala Leu Pro
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Pro Pro Leu Ser Met Pro Ser Thr Ser His Cys Thr Pro Pro Pro
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Tyr Pro Thr Asp Cys Ser Ile Val Ser Phe Leu Ala Arg Leu Gly Cys
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 Leu
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 Val
 Phe

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 His
 Ile
 Trp
 Asp
 Phe
 Leu
 Glu
 Gln
 Pro
 Ile
 Cys
 Ser
 Val
 Gln
 Pro

 Ile
 Asp
 Leu
 Asp
 Glu
 Pro
 Ser
 Glu
 Asp
 Gly
 Ala
 Thr
 Asn

 Lys
 Ile
 Glu
 Ile
 Ser
 Met
 Asp
 Cys
 Ile
 Arg
 Met
 Asp
 Ser
 Asp
 Leu
 Asp
 Leu
 Asp
 Leu
 Asp
 Leu
 Asp
 Leu
 Asp
 Leu
 Asp
 Ile
 Asp
 Ile

				٥٠											
The	7 ~~	114 0	70.7	85	70	0	77 - 7	m).	90	Б		-	-	95	
Inr	ASP	HIS	100	GIN	Asn	Ser	Val	105	Ala	Pro	Ser	Pro	Tyr 110	Ala	GIn
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	130				Gly	135					140				
Ser 145	Ser	Thr	Ala	Lys	Ser 150	Ala	Thr	Trp	Thr	Tyr 155	Ser	Thr	Glu	Leu	Lys 160
Lys	Leu	Tyr	Суѕ	Gln 165	Ile	Ala	Lys	Thr	Cys 170	Pro	Ile	Gln	Ile	Lys 175	Val
Met	Thr	Pro	Pro 180	Pro	Gln	Gly	Ala	Val 185	Ile	Arg	Ala	Met	Pro 190	Val	Tyr
Lys	Lys	Ala 195	Glu	His	Val	Thr	Glu 200	Val	Val	Lys	Arg	Cys 205	Pro	Asn	His
Glu	Leu 210	Ser	Arg	Glu	Phe	Asn 215	Glu	Gly	Gln	Ile	Ala 220	Pro	Pro	Ser	His
Leu 225	Ile	Arg	Val	Glu	Gly 230	Asn	Ser	His	Ala	Gln 235	Tyr	Val	Glu	Asp	Pro 240
Ile	Thr	Gly	Arg	Gln 245	Ser	Val	Leu	Val	Pro 250	Tyr	Glu	Pro	Pro	Gln 255	Val
Gly	Thr	Glu	Phe 260	Thr	Thr	Val	Leu	Tyr 265	Asn	Phe	Met	Cys	Asn 270	Ser	Ser
		275			Asn		280					285			
	290				Gln	295					300				_
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				325	Ser				330					335	_
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		355			Asp		360					365		_	_
	370				Met	375					380				
385					Gln 390					395					400
				405	His				410					415	
			420		Asn			425					430		
		435			Ser		440					445			
	450				Thr	455					460				
Pro 465	Met	Met	Gly	Thr	His 470	Met	Pro	Met	Ala	Gly 475	Asp	Met	Asn	Gly	Leu 480
				485	Leu				490					495	
			500		Pro			505					510		
Phe	Leu	Ala	Arg	Leu	Gly	Cys	Ser	Ser	Cys	Leu	Asp	Tyr	Phe	Thr	Thr

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545 550 555 560
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Leu Leu Arg Thr Pro Ser Ser Ala Ser Thr Val Ser Val Gly Ser Ser
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Gln Thr Ile Ser Phe Pro Pro Arg Asp Glu Trp Asn Asp Phe Asn Phe
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Lys Ile Glu Ile Ser Met Asp Cys Ile Arg Met Gln Asp Ser Asp Leu
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Ser Asp Pro Met Trp Pro Gln Tyr Thr Asn Leu Gly Leu Leu Asn Ser
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Met Asp Gln Gln Ile Gln Asn Gly Ser Ser Ser Thr Ser Pro Tyr Asn
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Thr Asp His Ala Gln Asn Ser Val Thr Ala Pro Ser Pro Tyr Ala Gln
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Pro Ser Ser Thr Phe Asp Ala Leu Ser Pro Ser Pro Ala Ile Pro Ser
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Asn Thr Asp Tyr Pro Gly Pro His Ser Phe Asp Val Ser Phe Gln Gln
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Ser Ser Thr Ala Lys Ser Ala Thr Trp Thr Tyr Ser Thr Glu Leu Lys
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Lys Leu Tyr Cys Gln Ile Ala Lys Thr Cys Pro Ile Gln Ile Lys Val
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Met Thr Pro Pro Pro Gln Gly Ala Val Ile Arg Ala Met Pro Val Tyr
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Lys Lys Ala Glu His Val Thr Glu Val Val Lys Arg Cys Pro Asn His
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Glu Leu Ser Arg Glu Phe Asn Glu Gly Gln Ile Ala Pro Pro Ser His
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Leu Ile Arg Val Glu Gly Asn Ser His Ala Gln Tyr Val Glu Asp Pro
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115

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Ile Thr Gly Arg Gln Ser Val Leu Val Pro Tyr Glu Pro Pro Gln Val
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Gly Thr Glu Phe Thr Thr Val Leu Tyr Asn Phe Met Cys Asn Ser Ser
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Cys Val Gly Gly Met Asn Arg Arg Pro Ile Leu Ile Ile Val Thr Leu
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Glu Thr Arg Asp Gly Gln Val Leu Gly Arg Arg Cys Phe Glu Ala Arg
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                                         300
Ile Cys Ala Cys Pro Gly Arg Asp Arg Lys Ala Asp Glu Asp Ser Ile
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Arg Lys Gln Gln Val Ser Asp Ser Thr Lys Asn Gly Asp Gly Thr Lys
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Arg Pro Phe Arg Gln Asn Thr His Gly Ile Gln Met Thr Ser Ile Lys
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Lys Arg Arg Ser Pro Asp Asp Glu Leu Leu Tyr Leu Pro Val Arg Gly
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Arg Glu Thr Tyr Glu Met Leu Leu Lys Ile Lys Glu Ser Leu Glu Leu
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                             380
Met Gln Tyr Leu Pro Gln His Thr Ile Glu Thr Tyr Arg Gln Gln Gln
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Gln Gln Gln His Gln His Leu Leu Gln Lys His Leu Leu Ser Ala Cys
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Val Thr Ala Pro Ser Pro Tyr Ala Gln Pro Ser Ser Thr Phe Asp Ala
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Leu Ser Pro Ser Pro Ala Ile Pro Ser Asn Thr Asp Tyr Pro Gly Pro
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                                     75
His Ser Phe Asp Val Ser Phe Gln Gln Ser Ser Thr Ala Lys Ser Ala
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Thr Trp Thr Tyr Ser Thr Glu Leu Lys Lys Leu Tyr Cys Gln Ile Ala
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                             105
Lys Thr Cys Pro Ile Gln Ile Lys Val Met Thr Pro Pro Pro Gln Gly
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Glu Val Val Lys Arg Cys Pro Asn His Glu Leu Ser Arg Glu Phe Asn

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Leu Val Pro Tyr Glu Pro Pro Gln Val Gly Thr Glu Phe Thr Thr Val
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Leu Tyr Asn Phe Met Cys Asn Ser Ser Cys Val Gly Gly Met Asn Arg
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                                         220
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Leu Gly Arg Arg Cys Phe Glu Ala Arg Ile Cys Ala Cys Pro Gly Arg
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His Gly Ile Gln Met Thr Ser Ile Lys Lys Arg Arg Ser Pro Asp Asp
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Arg Met Gln Asp Ser Asp Leu Ser Asp Pro Met Trp Pro Gln Tyr Thr
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                             105
Asn Leu Gly Leu Leu Asn Ser Met Asp Gln Gln Ile Gln Asn Gly Ser
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Ser Ser Thr Ser Pro Tyr Asn Thr Asp His Ala Gln Asn Ser Val Thr
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                                         140
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Ala Pro Ser Pro Tyr Ala Gln Pro Ser Ser Thr Phe Asp Ala Leu Ser

1 4 5					1 - 0										
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PIO	ser	PIO	Ald	165	Pro	ser	Asn	Thr	170	Tyr	Pro	GLY	Pro	His	Ser
Phe	Asn	Val	Ser		Gln	Gln	Sar	Sar		7\1 >	Tuc	Sor	717	175 Thr	Trn
1110	1100	Val	180	1110	0.111	GIII	per	185	1111	Ата	пуз	ser	190	1111	пр
Thr	Tvr	Ser		Glu	Leu	Lvs	Lvs		Tur	Cvs	Gln	Tle		Lys	Thr
	- 1	195					200		- 1 -	010	0111	205	111.0	БуО	1111
Cys	Pro	Ile	Gln	Ile	Lys	Val		Thr	Pro	Pro	Pro		Glv	Ala	Val
	210				_	215					220		_		
Ile	Arg	Ala	Met	Pro	Val	Tyr	Lys	Lys	Ala	Glu	His	Val	Thr	Glu	Val
225					230					235					240
Val	Lys	Arg	Cys		Asn	His	Glu	Leu		Arg	Glu	Phe	Asn	Glu	Gly
C1	T 1 -	71 T .	Б	245	0		_	- 1	250				_	255	
GTII	тте	Ата	260	Pro	ser	HlS	Leu		Arg	Val	Glu	GLY		Ser	His
ΔΙα	Gln	Тиг		Glu	7) an	Dro	TIO	265	Clar	7) ~~ ~	Cl.	Co~	270	Leu	17.0.7
1110	OIII	275	val	OIU	лэр	110	280	1111.	дту	Arg	GIII	285	Val	ьеи	val
Pro	Tvr		Pro	Pro	Gln	Val		Thr	Glu	Phe	Thr		Val	Leu	Tur
	290					295	1		0_0		300			Lea	- y -
Asn	Phe	Met	Cys	Asn	Ser	Ser	Cys	Val	Gly	Gly		Asn	Arq	Arg	Pro
305					310					315					320
Ile	Leu	Ile	Ile		Thr	Leu	Glu	Thr	Arg	Asp	Gly	Gln	Val	Leu	Gly
_	_	_		325	_				330					335	
Arg	Arg	Суѕ		Glu	Ala	Arg	Ile		Ala	Cys	Pro	Gly		Asp	Arg
Tric	7.7.5	7 000	340	71 ~~~	C	т1.	7	345	C1.	01	17.7	~	350	_	m.
ьуѕ	MId	355	GIU	ASP	ser	тте	360	гуѕ	GIN	GIN	vaı	365	Asp	Ser	Tnr
Lvs	Asn		Asp	Glv	Thr	Lvs		Pro	Phe	Ara	Gln		Thr	His	Glv
	370	1		1		375	9		1110	9	380	11011	1111	1110	O L y
Ile	Gln	Met	Thr	Ser	Ile	Lys	Lys	Arg	Arg	Ser	Pro	Asp	Asp	Glu	Leu
385					390					395					400
Leu	Tyr	Leu	Pro		Arg	Gly	Arg	Glu		Tyr	Glu	Met	Leu	Leu	Lys
T1.	T	C 1	0	405	~ 3	-		~ 1	410	_			•	415	
тте	ьуs	GIU	Ser 420	Leu	GLu	Leu	Met		Tyr	Leu	Pro	Gln		Thr	Ile
Glu	Thr	Tur		Gln	Gla	Cln	Cln	425	Cln	uic	Cln	шіс	430	Leu	C1~
OIU	1111	435	Arg	Gill	GIII	G1.11	440	GIII	GIII	пто	GIII	445	ьeu	ьеи	GIII
Lys	Gln		Ser	Ile	Gln	Ser		Ser	Ser	Tvr	Glv	-	Ser	Ser	Pro
-	450					455				- 1 -	460		~ ~ _		
Pro	Leu	Asn	Lys	Met	Asn	Ser	Met	Asn	Lys	Leu	Pro	Ser	Val	Ser	Gln
465					470					475					480
Leu	Ile	Asn	Pro		Gln	Arg	Asn	Ala	Leu	Thr	Pro	Thr	Thr	Ile	Pro
	<i>a</i> 1		~ 1	485	_		_		490					495	
Asp	GTA	Met		Ala	Asn	Ile	Pro		Met	Gly	Thr	His		Pro	Met
ΔΊο	Clv	Aen	500	Acn	C111	T 011	Cor	505	Π.b.∞	C1 ~	7.3.0	T 0	510	Desa	D
ALG	Сту	515	nec	MSII	Gry	пеп	520	PLO	1111	GIII	Ald	ьеи 525	Pro	Pro	Pro
Leu	Ser		Pro	Ser	Thr	Ser		Cvs	Thr	Pro	Pro		Pro	Tyr	Pro
	530					535		- 10	~ ~ ~ ~		540			- J -	110
Thr		Cys	Ser	Ile	Val		Phe	Leu	Ala	Arq		Gly	Cys	Ser	Ser
545					550					555					560
Cys	Leu	Asp	Tyr	Phe	Thr	Thr	Gln	Gly	Leu	Thr	Thr	Ile	Tyr	Gln	
				565					570					575	
Glu	His	Tyr	Ser	Met	Asp	Asp	Leu	Ala	Ser	Leu	Lys	Ile	Pro	Glu	Gln

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580
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Phe Arg His Ala Ile Trp Lys Gly Ile Leu Asp His Arg Gln Leu His
      595 600 605
Glu Phe Ser Ser Pro Ser His Leu Leu Arg Thr Pro Ser Ser Ala Ser
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                                       620
Thr Val Ser Val Gly Ser Ser Glu Thr Arg Gly Glu Arg Val Ile Asp
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                      635
Ala Val Arg Phe Thr Leu Arg Gln Thr Ile Ser Phe Pro Pro Arg Asp
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Glu Trp Asn Asp Phe Asn Phe Asp Met Asp Ala Arg Arg Asn Lys Gln
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Gln Arg Ile Lys Glu Glu Gly Glu
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Tyr Thr Asn Leu Gly Leu Leu Asn Ser Met Asp Gln Gln Ile Gln Asn
                             25
Gly Ser Ser Ser Thr Ser Pro Tyr Asn Thr Asp His Ala Gln Asn Ser
                         4.0
Val Thr Ala Pro Ser Pro Tyr Ala Gln Pro Ser Ser Thr Phe Asp Ala
                  55
Leu Ser Pro Ser Pro Ala Ile Pro Ser Asn Thr Asp Tyr Pro Gly Pro
                 70
                                    75
His Ser Phe Asp Val Ser Phe Gln Gln Ser Ser Thr Ala Lys Ser Ala
             85
                                90
Thr Trp Thr Tyr Ser Thr Glu Leu Lys Lys Leu Tyr Cys Gln Ile Ala
          100
                            105
Lys Thr Cys Pro Ile Gln Ile Lys Val Met Thr Pro Pro Pro Gln Gly
                         120
                                           125
Ala Val Ile Arg Ala Met Pro Val Tyr Lys Lys Ala Glu His Val Thr
                     135
                                       140
Glu Val Val Lys Arg Cys Pro Asn His Glu Leu Ser Arg Glu Phe Asn
                  150
                                    155
Glu Gly Gln Ile Ala Pro Pro Ser His Leu Ile Arg Val Glu Gly Asn
              165
                                170
Ser His Ala Gln Tyr Val Glu Asp Pro Ile Thr Gly Arg Gln Ser Val
                             185
                                               190
Leu Val Pro Tyr Glu Pro Pro Gln Val Gly Thr Glu Phe Thr Thr Val
      195
                         200
                                           205
Leu Tyr Asn Phe Met Cys Asn Ser Ser Cys Val Gly Gly Met Asn Arg
                    215
                                       220
Arg Pro Ile Leu Ile Ile Val Thr Leu Glu Thr Arg Asp Gly Gln Val
     230
                       235
Leu Gly Arg Arg Cys Phe Glu Ala Arg Ile Cys Ala Cys Pro Gly Arg
              245
                                250
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Asp Arg Lys Ala Asp Glu Asp Ser Ile Arg Lys Gln Gln Val Ser Asp

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260
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                                          270
Ser Thr Lys Asn Gly Asp Gly Thr Lys Arg Pro Phe Arg Gln Asn Thr
            280 285
His Gly Ile Gln Met Thr Ser Ile Lys Lys Arg Arg Ser Pro Asp Asp
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                                    300
Glu Leu Leu Tyr Leu Pro Val Arg Gly Arg Glu Thr Tyr Glu Met Leu
305 310 315
Leu Lys Ile Lys Glu Ser Leu Glu Leu Met Gln Tyr Leu Pro Gln His
            325
                             330
Thr Ile Glu Thr Tyr Arg Gln Gln Gln Gln Gln His Gln His Leu
          340
                          345
Leu Gln Lys Gln Thr Ser Ile Gln Ser Pro Ser Ser Tyr Gly Asn Ser
                       360
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Ser Pro Pro Leu Asn Lys Met Asn Ser Met Asn Lys Leu Pro Ser Val
                          380
                   375
Ser Gln Leu Ile Asn Pro Gln Gln Arg Asn Ala Leu Thr Pro Thr Thr
                390
                       395
Ile Pro Asp Gly Met Gly Ala Asn Ile Pro Met Met Gly Thr His Met
            405
                 410 415
Pro Met Ala Gly Asp Met Asn Gly Leu Ser Pro Thr Gln Ala Leu Pro
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Pro Pro Leu Ser Met Pro Ser Thr Ser His Cys Thr Pro Pro Pro
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Tyr Pro Thr Asp Cys Ser Ile Val Arg Ile Trp Gln Val
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                          25
Ile Asp Leu Asn Phe Val Asp Glu Pro Ser Glu Asp Gly Ala Thr Asn
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Lys Ile Glu Ile Ser Met Asp Cys Ile Arg Met Gln Asp Ser Asp Leu
                    55
Ser Asp Pro Met Trp Pro Gln Tyr Thr Asn Leu Gly Leu Leu Asn Ser
                70
                                75
Met Asp Gln Gln Ile Gln Asn Gly Ser Ser Ser Thr Ser Pro Tyr Asn
                              90
Thr Asp His Ala Gln Asn Ser Val Thr Ala Pro Ser Pro Tyr Ala Gln
                         105
                                           110
Pro Ser Ser Thr Phe Asp Ala Leu Ser Pro Ser Pro Ala Ile Pro Ser
                            125
                       120
Asn Thr Asp Tyr Pro Gly Pro His Ser Phe Asp Val Ser Phe Gln Gln
 130 135 140
Ser Ser Thr Ala Lys Ser Ala Thr Trp Thr Tyr Ser Thr Glu Leu Lys
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150 155 160

Lys Leu Tyr Cys Gln Ile Ala Lys Thr Cys Pro Ile Gln Ile Lys Val

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Met Thr Pro Pro Pro Gln Gly Ala Val Ile Arg Ala Met Pro Val Tyr
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Lys Lys Ala Glu His Val Thr Glu Val Val Lys Arg Cys Pro Asn His
           200
Glu Leu Ser Arg Glu Phe Asn Glu Gly Gln Ile Ala Pro Pro Ser His
                     215
Leu Ile Arg Val Glu Gly Asn Ser His Ala Gln Tyr Val Glu Asp Pro
                 230
                                   235
Ile Thr Gly Arg Gln Ser Val Leu Val Pro Tyr Glu Pro Pro Gln Val
              245
                                250
Gly Thr Glu Phe Thr Thr Val Leu Tyr Asn Phe Met Cys Asn Ser Ser
          260
                            265
Cys Val Gly Met Asn Arg Arg Pro Ile Leu Ile Ile Val Thr Leu
                        280
Glu Thr Arg Asp Gly Gln Val Leu Gly Arg Arg Cys Phe Glu Ala Arg
                     295
                                      300
Ile Cys Ala Cys Pro Gly Arg Asp Arg Lys Ala Asp Glu Asp Ser Ile
                310
                      315
Arg Lys Gln Gln Val Ser Asp Ser Thr Lys Asn Gly Asp Gly Thr Lys
             325 330
Arg Pro Phe Arg Gln Asn Thr His Gly Ile Gln Met Thr Ser Ile Lys
                            345
Lys Arg Arg Ser Pro Asp Asp Glu Leu Leu Tyr Leu Pro Val Arg Gly
       355
                         360
Arg Glu Thr Tyr Glu Met Leu Leu Lys Ile Lys Glu Ser Leu Glu Leu
                     375
Met Gln Tyr Leu Pro Gln His Thr Ile Glu Thr Tyr Arg Gln Gln Gln
                 390
                                   395
Gln Gln His Gln His Leu Leu Gln Lys Gln Thr Ser Ile Gln Ser
             405
                            410
Pro Ser Ser Tyr Gly Asn Ser Ser Pro Pro Leu Asn Lys Met Asn Ser
                         425 430
Met Asn Lys Leu Pro Ser Val Ser Gln Leu Ile Asn Pro Gln Gln Arg
                        440 445
      435
Asn Ala Leu Thr Pro Thr Thr Ile Pro Asp Gly Met Gly Ala Asn Ile
                     455
                                       460
Pro Met Met Gly Thr His Met Pro Met Ala Gly Asp Met Asn Gly Leu
                 470
                                    475
Ser Pro Thr Gln Ala Leu Pro Pro Pro Leu Ser Met Pro Ser Thr Ser
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His Cys Thr Pro Pro Pro Pro Tyr Pro Thr Asp Cys Ser Ile Val Arg
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Ile Trp Gln Val
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<212> DNA

<213> Homo sapiens

<400> 345

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tgacattcgt atcatcactg tgcaccattg gcttctaggc actccagtgg ggtaggagaa 180
ggaggtetga aaccetegea gagggatett geceteatte tittgggtetg aaacaetgge 240
agtegttgga aacaggacte agggataaac cagegeaatg gattggggga egetgeacac 300
tttcatcggg ggtgtcaaca aacactccac cagcatcggg aaggtgtgga tcacagtcat 360
ctttattttc cgagtcatga tcctagtggt ggctgcccag gaagtgtggg gtgacgagca 420
agaggactic gtctgcaaca cactgcaacc gggatgcaaa aatgtgtgct atgaccactt 480
tttcccggtg tcccacatcc ggctgtgggc cctccagctg atcttcgtct ccaccccage 540
gctgctggtg gccatgcatg tggcctacta caggcacgaa accactcgca agttcaggcg 600
aggagagaag aggaatgatt tcaaagacat agaggacatt aaaaagcaca aggttcggat 660
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agoctttatg tatgtgtttt actteettta eaatgggtae eacetgeeet gggtgttgaa 780
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gaccgtgttt accattttta tgatttctgc gtctgtgatt tgcatgctgc ttaacgtggc 900
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ttcagatagt ggtcaaaatg caatcacagg tttcccaagc taaacatttc aaggtaaaat 1080
gtagctgcgt cataaggaga cttctgtctt ctccagaagg caataccaac ctgaaagttc 1140
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gactototga caaagtgggt actttotgaa aatttatata actgttgttg ataaggaaca 1380
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tacgcttaag gtgggaaagt gttcattgca caatatattt ttactgcttt ctgaatgtag 1680
acqgaacaqt qtqqaaqcaq aaqqcttttt taactcatcc qtttqqccqa tcqttqcaqa 1740
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Ser Thr Ser Ile Gly Lys Val Trp Ile Thr Val Ile Phe Ile Phe Arg
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                                                   30
Val Met Ile Leu Val Val Ala Ala Gln Glu Val Trp Gly Asp Glu Gln
                           40
                                               45
Glu Asp Phe Val Cys Asn Thr Leu Gln Pro Gly Cys Lys Asn Val Cys
                        55
Tyr Asp His Phe Phe Pro Val Ser His Ile Arg Leu Trp Ala Leu Gln
                    70
                                       75
Leu Ile Phe Val Ser Thr Pro Ala Leu Leu Val Ala Met His Val Ala
                85
                                   90
Tyr Tyr Arg His Glu Thr Thr Arg Lys Phe Arg Arg Gly Glu Lys Arg
            100
                               105
Asn Asp Phe Lys Asp Ile Glu Asp Ile Lys Lys His Lys Val Arg Ile
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                                               125
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Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser Ser Ile Phe Phe Arg Ile

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140
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Ile Phe Glu Ala Ala Phe Met Tyr Val Phe Tyr Phe Leu Tyr Asn Gly
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Tyr His Leu Pro Trp Val Leu Lys Cys Gly Ile Asp Pro Cys Pro Asn
                                    170
                165
Leu Val Asp Cys Phe Ile Ser Arg Pro Thr Glu Lys Thr Val Phe Thr
                                185
                                                    190
            180
Ile Phe Met Ile Ser Ala Ser Val Ile Cys Met Leu Leu Asn Val Ala
                            200
Glu Leu Cys Tyr Leu Leu Leu Lys Val Cys Phe Arg Arg Ser Lys Arg
                        215
                                            220
Ala Gln Thr Gln Lys Asn His Pro Asn His Ala Leu Lys Glu Ser Lys
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                                        235
Gln Asn Glu Met Asn Glu Leu Ile Ser Asp Ser Gly Gln Asn Ala Ile
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Thr Gly Phe Pro Ser
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ttegtggaet geeeggaega gagetgggee eteaaggeea tegaggeget tteaggtaaa 180
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cqqaaacttc agatacqaaa tatcccgcct catttacagt gggaggtgct ggatagttta 300
ctagtccagt atggagtggt ggagagctgt gagcaagtga acactgactc ggaaactgca 360
qttqtaaatq taacctattc caqtaaqqac caaqctaqac aaqcactaqa caaactqaat 420
ggatttcagt tagagaattt caccttgaaa gtagcctata tccctgatga aacggccgcc 480
cagcaaaacc cettgcagca geecegaggt egeeggggge ttgggcagag gggeteetea 540
aggcaggggt ctccaggatc cgtatccaag cagaaaccat gtgatttgcc tctgcgcctg 600
ctggttccca cccaatttgt tggagccatc ataggaaaag aaggtgccac cattcggaac 660
atcaccaaac agacccagte taaaatcgat gtccaccgta aagaaaatgc gggggctgct 720
gagaagtega ttactateet etetaeteet gaaggeacet etgeggettg taagtetatt 780
ctggagatta tgcataagga agctcaagat ataaaattca cagaagagat ccccttgaag 840
attttagctc ataataactt tgttggacgt cttattggta aagaaggaag aaatcttaaa 900
aaaattgagc aagacacaga cactaaaatc acgatatctc cattgcagga attgacgctg 960
tataatccag aacgcactat tacagttaaa ggcaatgttg agacatgtgc caaagctgag 1020
gaggagatca tgaagaaaat cagggagtct tatgaaaatg atattgcttc tatgaatctt 1080
caagcacatt taatteetgg attaaatetg aacgeettgg gtetgtteee acceaettea 1140
gggatgecae eteccacete agggeeecet teagecatga etecteceta ecegeagttt 1200
gagcaatcag aaacggagac tgttcatctg tttatcccag ctctatcagt cggtgccatc 1260
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attgctccag cggaagcacc agatgctaaa gtgaggatgg tgattatcac tggaccacca 1380
gaggeteagt teaaggetea gggaagaatt tatggaaaaa ttaaagaaga aaaetttgtt 1440
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gttqttqtcc ctcqtqacca qacacctqat qaqaatqacc aagtqqttqt caaaataact 1620
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<212> PRT
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Phe Leu Val Lys Thr Gly Tyr Ala Phe Val Asp Cys Pro Asp Glu Ser
                                     45
                         40
Trp Ala Leu Lys Ala Ile Glu Ala Leu Ser Gly Lys Ile Glu Leu His
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Gly Lys Pro Ile Glu Val Glu His Ser Val Pro Lys Arg Gln Arg Ile
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Arg Lys Leu Gln Ile Arg Asn Ile Pro Pro His Leu Gln Trp Glu Val
              85
                                90
Leu Asp Ser Leu Leu Val Gln Tyr Gly Val Val Glu Ser Cys Glu Gln
 100 105
Val Asn Thr Asp Ser Glu Thr Ala Val Val Asn Val Thr Tyr Ser Ser
                         120
                                            125
Lys Asp Gln Ala Arg Gln Ala Leu Asp Lys Leu Asn Gly Phe Gln Leu
                      135
                                        140
Glu Asn Phe Thr Leu Lys Val Ala Tyr Ile Pro Asp Glu Thr Ala Ala
                                    155
                 150
Gln Gln Asn Pro Leu Gln Gln Pro Arg Gly Arg Arg Gly Leu Gly Gln
                                170
Arg Gly Ser Ser Arg Gln Gly Ser Pro Gly Ser Val Ser Lys Gln Lys
                            185
          180
                                               190
Pro Cys Asp Leu Pro Leu Arg Leu Leu Val Pro Thr Gln Phe Val Gly
       195 200 205
Ala Ile Ile Gly Lys Glu Gly Ala Thr Ile Arg Asn Ile Thr Lys Gln
 210 215 220
Thr Gln Ser Lys Ile Asp Val His Arg Lys Glu Asn Ala Gly Ala Ala
                 230
                                    235
Glu Lys Ser Ile Thr Ile Leu Ser Thr Pro Glu Gly Thr Ser Ala Ala
              245
                                 250
Cys Lys Ser Ile Leu Glu Ile Met His Lys Glu Ala Gln Asp Ile Lys
                             265
Phe Thr Glu Glu Ile Pro Leu Lys Ile Leu Ala His Asn Asn Phe Val
                        280
Gly Arg Leu Ile Gly Lys Glu Gly Arg Asn Leu Lys Lys Ile Glu Gln
                      295
                                        300
Asp Thr Asp Thr Lys Ile Thr Ile Ser Pro Leu Gln Glu Leu Thr Leu
                 310
                                    315
Tyr Asn Pro Glu Arg Thr Ile Thr Val Lys Gly Asn Val Glu Thr Cys
              325
                                 330
Ala Lys Ala Glu Glu Glu Ile Met Lys Lys Ile Arq Glu Ser Tyr Glu
                            345
Asn Asp Ile Ala Ser Met Asn Leu Gln Ala His Leu Ile Pro Gly Leu
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                                            365
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Asn Leu Asn Ala Leu Gly Leu Phe Pro Pro Thr Ser Gly Met Pro Pro
                        375
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Pro Thr Ser Gly Pro Pro Ser Ala Met Thr Pro Pro Tyr Pro Gln Phe
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                                        395
Glu Gln Ser Glu Thr Glu Thr Val His Leu Phe Ile Pro Ala Leu Ser
                405
                                    410
                                                         415
Val Gly Ala Ile Ile Gly Lys Gln Gly Gln His Ile Lys Gln Leu Ser
                                425
Arg Phe Ala Gly Ala Ser Ile Lys Ile Ala Pro Ala Glu Ala Pro Asp
                            440
                                                 445
        435
Ala Lys Val Arq Met Val Ile Ile Thr Gly Pro Pro Glu Ala Gln Phe
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                        455
                                             460
Lys Ala Gln Gly Arg Ile Tyr Gly Lys Ile Lys Glu Glu Asn Phe Val
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                                        475
Ser Pro Lys Glu Glu Val Lys Leu Glu Ala His Ile Arg Val Pro Ser
                485
                                    490
Phe Ala Ala Gly Arg Val Ile Gly Lys Gly Gly Lys Thr Val Asn Glu
            500
                                505
                                                     510
Leu Gln Asn Leu Ser Ser Ala Glu Val Val Val Pro Arg Asp Gln Thr
        515
                            520
                                                 525
Pro Asp Glu Asn Asp Gln Val Val Lys Ile Thr Gly His Phe Tyr
                        535
    530
                                             540
Ala Cys Gln Val Ala Gln Arg Lys Ile Gln Glu Ile Leu Thr Gln Val
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Lys Gln His Gln Gln Gln Lys Ala Leu Gln Ser Gly Pro Pro Gln Ser
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Arg Arg Lys
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gaaaagatga gagaagttac agacteteet gggegaeeee gagagettae eatteeteag 180
acttcttcac atggtgctaa cagattt
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Ser Ser Gln Ile Ala Ala Ala Ser Thr Gln Pro Glu Asp Asp Ile
            20
                                25
                                                     30
Asn Thr Gln Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val Thr Asp
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Ser Pro Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr Ser Ser His

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<210> 351
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ccqatcqqqc aqqcqatqqc qatcqcqqqc caqatcaaqc ttcccaccqt tcatatcqqq 180
cctaccgcct tcctcggctt gggtgttgtc gacaacaacg gcaacggcgc acgagtccaa 240
egeqtqqteq qqaqeqetee qqeqqeaaqt eteqqeatet ceaeeqqeqa egtqateaee 300
geggtegacg gegeteegat caacteggee accgegatgg eggacgeget taaegggeat 360
cateceggtg acgtcatete ggtgacetgg caaaccaagt egggeggeae gegtacaggg 420
aacgtgacat tggccgaggg acccccggcc gaattcatgg attgggggac gctgcacact 480
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gaggaetteg tetgeaacae actgeaaceg ggatgeaaaa atgtgtgeta tgaceaettt 660
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ggagagaaga ggaatgattt caaagacata gaggacatta aaaagcagaa ggttcggata 840
gaggggtgac tegageacca ecaecaceae caetgagate eggetgetaa caaageeega 900
aaggaagetg agttggetge tgecaceget gagcaataac tagcataace cettggggee 960
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Ser Gln Gly Gln Gly Phe Ala Ile Pro Ile Gly Gln Ala Met Ala
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Ile Ala Gly Gln Ile Lys Leu Pro Thr Val His Ile Gly Pro Thr Ala
Phe Leu Gly Leu Gly Val Val Asp Asn Asn Gly Asn Gly Ala Arg Val
                        55
                                            60
Gln Arg Val Val Gly Ser Ala Pro Ala Ala Ser Leu Gly Ile Ser Thr
                    70
                                        75
Gly Asp Val Ile Thr Ala Val Asp Gly Ala Pro Ile Asn Ser Ala Thr
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Ala Met Ala Asp Ala Leu Asn Gly His His Pro Gly Asp Val Ile Ser
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Val Thr Trp Gln Thr Lys Ser Gly Gly Thr Arg Thr Gly Asn Val Thr
        115
                            120
                                                125
Leu Ala Glu Gly Pro Pro Ala Glu Phe Met Asp Trp Gly Thr Leu His
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                                            140
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Thr Phe Ile Gly Gly Val Asn Lys His Ser Thr Ser Ile Gly Lys Val

50

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150
145
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Trp Ile Thr Val Ile Phe Ile Phe Arg Val Met Ile Leu Val Val Ala
                165
                                    170
Ala Gln Glu Val Trp Gly Asp Glu Gln Glu Asp Phe Val Cys Asn Thr
            180
                                185
Leu Gln Pro Gly Cys Lys Asn Val Cys Tyr Asp His Phe Pro Val
                            200
                                                205
Ser His Ile Arg Leu Trp Ala Leu Gln Leu Ile Phe Val Ser Thr Pro
                        215
Ala Leu Leu Val Ala Met His Val Ala Tyr Tyr Arg His Glu Thr Thr
                    230
                                        235
Arg Lys Phe Arg Arg Gly Glu Lys Arg Asn Asp Phe Lys Asp Ile Glu
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Asp Ile Lys Lys Gln Lys Val Arg Ile Glu Gly
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accepticata tegggeetae egeetteete ggettgggtg tigtegaeaa eaacggeaae 180
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aaqaqaqcac aqacqcaaaa aaatcacccc aatcatqccc taaaqqaqaq taaqcaqaat 840
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Ile Ala Gly Gln Ile Lys Leu Pro Thr Val His Ile Gly Pro Thr Ala
                            40
Phe Leu Gly Leu Gly Val Val Asp Asn Asn Gly Asn Gly Ala Arg Val
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55

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Gln Arg Val Val Gly Ser Ala Pro Ala Ala Ser Leu Gly Ile Ser Thr
Gly Asp Val Ile Thr Ala Val Asp Gly Ala Pro Ile Asn Ser Ala Thr
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Ala Met Ala Asp Ala Leu Asn Gly His His Pro Gly Asp Val Ile Ser
        100 105
Val Thr Trp Gln Thr Lys Ser Gly Gly Thr Arg Thr Gly Asn Val Thr
      115 120
                            125
Leu Ala Glu Gly Pro Pro Ala Glu Phe His Glu Thr Thr Arg Lys Phe
                        140
                   135
Arg Arg Gly Glu Lys Arg Asn Asp Phe Lys Asp Ile Glu Asp Ile Lys
                    155
     150
Lys Gln Lys Val Arg Ile Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser
            165
                   170 175
Ser Ile Phe Phe Arq Ile Ile Phe Glu Ala Ala Phe Met Tyr Val Phe
        180 185 190
Tyr Phe Leu Tyr Asn Gly Tyr His Leu Pro Trp Val Leu Lys Cys Gly
 195 200
                                      205
Ile Asp Pro Cys Pro Asn Leu Val Asp Cys Phe Ile Ser Arg Pro Thr
                   215 220
Glu Lys Thr Val Phe Thr Ile Phe Met Ile Ser Ala Ser Val Ile Cys
   230
                                235 240
Met Leu Leu Asn Val Ala Glu Leu Cys Tyr Leu Leu Leu Lys Val Cys
            245
                             250
Phe Arg Arg Ser Lys Arg Ala Gln Thr Gln Lys Asn His Pro Asn His
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Ser Gly Gln Asn Ala Ile Thr Gly Phe Pro Ser
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Asn Leu Ile Ser Asn Ile Lys Glu Met Ile Thr Glu Ala Ser Phe Tyr
Leu Phe Asn Ala Thr Lys Arg Arg Val Phe Phe Arg Asn Ile Lys Ile
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Leu Ile Pro Ala Thr Trp Lys Ala Asn Asn Ser Lys Ile Lys Gln
Glu Ser Tyr Glu Lys Ala Asn Val Ile Val Thr Asp Trp Tyr Gly Ala
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             8.5
His Gly Asp Asp Pro Tyr Thr Leu Gln Tyr Arg Gly Cys Gly Lys Glu
 100
                            105
                                               110
Gly Lys Tyr Ile His Phe Thr Pro Asn Phe Leu Leu Asn Asp Asn Leu
                               125
                         120
Thr Ala Gly Tyr Gly Ser Arg Gly Arg Val Phe Val His Glu Trp Ala
                     135
                                       140
His Leu Arg Trp Gly Val Phe Asp Glu Tyr Asn Asn Asp Lys Pro Phe
                  150
                                    155
Tyr Ile Asn Gly Gln Asn Gln Ile Lys Val Thr Arg Cys Ser Ser Asp
                                170
              165
Ile Thr Gly Ile Phe Val Cys Glu Lys Gly Pro Cys Pro Gln Glu Asn
         180 185
Cys Ile Ile Ser Lys Leu Phe Lys Glu Gly Cys Thr Phe Ile Tyr Asn
      195
                     200
                                           205
Ser Thr Gln Asn Ala Thr Ala Ser Ile Met Phe Met Gln Ser Leu Ser
                     215
                                       220
Ser Val Val Glu Phe Cys Asn Ala Ser Thr His Asn Gln Glu Ala Pro
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                      235
Asn Leu Gln Asn Gln Met Cys Ser Leu Arq Ser Ala Trp Asp Val Ile
                                250
Thr Asp Ser Ala Asp Phe His His Ser Phe Pro Met Asn Gly Thr Glu
           260
                             265
Leu Pro Pro Pro Pro Thr Phe Ser Leu Val Glu Ala Gly Asp Lys Val
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Val Cys Leu Val Leu Asp Val Ser Ser Lys Met Ala Glu Ala Asp Arg
                    295
                                       300
Leu Leu Gln Leu Gln Gln Ala Ala Glu Phe Tyr Leu Met Gln Ile Val
                 310
                                    315
Glu Ile His Thr Phe Val Gly Ile Ala Ser Phe Asp Ser Lys Gly Glu
              325
                                330
Ile Arg Ala Gln Leu His Gln Ile Asn Ser Asn Asp Asp Arg Lys Leu
          340
                             345
Leu Val Ser Tyr Leu Pro Thr Thr Val Ser Ala Lys Thr Asp Ile Ser
                         360
Ile Cys Ser Gly Leu Lys Lys Gly Phe Glu Val Val Glu Lys Leu Asn
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Gly 385	Lys	Ala	Tyr	Gly	Ser 390	Val	Met	Ile	Leu	Val 395	Thr	Ser	Gly	Asp	Asp 400
Lys	Leu	Leu	Gly	Asn 405	Cys	Leu	Pro	Thr	Val 410	Leu	Ser	Ser	Gly	Ser 415	Thr
Ile	His	Ser	Ile 420	Ala	Leu	Gly	Ser	Ser 425	Ala	Ala	Pro	Asn	Leu 430	Glu	Glu
Leu	Ser	Arg 435	Leu	Thr	Gly	Gly	Leu 440	Lys	Phe	Phe	Val	Pro 445	Asp	Ile	Ser
Asn	Ser 450	Asn	Ser	Met	Ile	Asp 455	Ala	Phe	Ser	Arg	Ile 460	Ser	Ser	Gly	Thr
Gly 465	Asp	Ile	Phe	Gln	Gln 470	His	Ile	Gln	Leu	Glu 475	Ser	Thr	Gly	Glu	Asn 480
Val	Lys	Pro	His	His 485	Gln	Leu	Lys	Asn	Thr 490	Val	Thr	Val	Asp	Asn 495	Thr
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Pro	Glu	Ile 515	Ile	Leu	Phe	Asp	Pro 520	Asp	Gly	Arg	Lys	Tyr 525	Tyr	Thr	Asn
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545					550		-		_	555				Thr	560
				565					570					Ser 575	
			580					585					590	Asp	
		595					600		_			605	_	Gln	_
	610					615					620			Pro	
625					630					635				Gly	640
				645		_		=	650	_	_			Ser 655	
			660					665					670	Ser	
		675					680			_		685		Met	
	690					695					700			Pro	
705					710					715					Arg 720
				725					730	_				Gly 735	
		_	740				_	745			_		750	Ala	
-		755					760		_			765	_	Glu	~
	770					775					780			Lys	
785					790					795					Thr 800
ser	ьуѕ	Arg	ASN	Pro 805	GIN	GIN	Ala	GLY	810	Arg	Glu	TTE	rhe	Thr 815	Pne

```
Ser Pro Gln Ile Ser Thr Asn Gly Pro Glu His Gln Pro Asn Gly Glu
                                825
Thr His Glu Ser His Arg Ile Tyr Val Ala Ile Arg Ala Met Asp Arg
                            840
                                                845
Asn Ser Leu Gln Ser Ala Val Ser Asn Ile Ala Gln Ala Pro Leu Phe
                        855
                                            860
Ile Pro Pro Asn Ser Asp Pro Val Pro Ala Arg Asp Tyr Leu Ile Leu
                    870
                                        875
Lys Gly Val Leu Thr Ala Met Gly Leu Ile Gly Ile Ile Cys Leu Ile
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atgatttatg ccaatgtgaa acagggattt tatcccattc ttaatgccac tgtcactgcc 1860
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acagttgage cagagactgg agatectgtt acgetgagae teettgatga tggageaggt 1920

gctgatgtta taaaaaatga tggaatttac tcgaggtatt ttttctcctt tgctgcaaat 1980 qqtagatata qcttgaaagt qcatgtcaat cactctccca qcataagcac cccaqcccac 2040

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     cgagtcagct caggaggete cttttcagtg ctgggagtte cagctggece ccaccctgat 2220
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ڊ
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Ξ
= F
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     Lys Met Arg Glu Val Thr Asp Ser Pro Gly Arg Pro Arg Glu Leu Thr
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     Ile Pro Gln Thr Ser Ser His Gly Ala Asn Arg Phe Val
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     aatacacaga ggaagaagag tcaggaaaag atgagagaag ttacagactc tcctgggcga 180
     eccegagage traceatree teagactret teacarggree craacagart regitted 240
     attc
     <210> 363
     <211> 20
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1 4 5
     <213> Homo sapiens
Street Turk Tool Tark
     <400> 363
     Met Trp Gln Pro Leu Phe Phe Lys Trp Leu Leu Ser Cys Cys Pro Gly
                                            10
     Ser Ser Gln Ile
* - B
                  20
۱. آ
پود
E.
:
     <210> 364
i ag
     <211> 60
n
     <212> DNA
dent
dent
     <213> Homo sapiens
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122
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gagcatette aacaatatte etacagttgt atgggtggca tgttggccat ttgtgatgta 2100 gccgaatata ggaagtgtgc caaagacttc aagattccaa tggtattaca tctttttgat 2160 actetgeatg etetttgeaa tettetggta gttgeeceag ataatttaaa geaagtetge 2220 tcaggagaac aacttgctaa tctggacaag aatatacttc actccttcgt acaacttcgt 2280 gctgattata gatctgcccg ccttgctcga cacttcagct gagattgaat ttacaaagga 2340 att <210> 369 <211> 708 <212> PRT <213> Homo sapiens <400> 369 Met Ala Thr Thr Ala Glu Leu Phe Glu Glu Pro Phe Val Ala Asp Glu 10 Tyr Ile Glu Arg Leu Val Trp Arg Thr Pro Gly Gly Ser Arg Gly 25 20 Gly Pro Glu Ala Phe Asp Pro Lys Arg Leu Leu Glu Glu Phe Val Asn 3.5 40 4.5 His Ile Gln Glu Leu Gln Ile Met Asp Glu Arg Ile Gln Arg Lys Val 55 60 Glu Lys Leu Glu Gln Gln Cys Gln Lys Glu Ala Lys Glu Phe Ala Lys 70 75 Lys Val Gln Glu Leu Gln Lys Ser Asn Gln Val Ala Phe Gln His Phe 85 90 Gln Glu Leu Asp Glu His Ile Ser Tyr Val Ala Thr Lys Val Cys His 100 105 110 Leu Gly Asp Gln Leu Glu Gly Val Asn Thr Pro Arg Gln Arg Ala Val 120 125 Glu Ala Gln Lys Leu Met Lys Tyr Phe Asn Glu Phe Leu Asp Gly Glu 135 140 Leu Lys Ser Asp Val Phe Thr Asn Ser Glu Lys Ile Lys Glu Ala Ala 150 155 Asp Ile Ile Gln Lys Leu His Leu Ile Ala Gln Glu Leu Pro Phe Asp 165 170 175 Arg Phe Ser Glu Val Lys Ser Lys Ile Ala Ser Lys Tyr His Asp Leu 185 Glu Cys Gln Leu Ile Gln Glu Phe Thr Ser Ala Gln Arg Arg Gly Glu 200 205 Ile Ser Arg Met Arg Glu Val Ala Ala Val Leu Leu His Phe Lys Gly 210 215 220 Tyr Ser His Cys Val Asp Val Tyr Ile Lys Gln Cys Gln Glu Gly Ala 230 235 Tyr Leu Arg Asn Asp Ile Phe Glu Asp Ala Gly Ile Leu Cys Gln Arg 245 250 Val Asn Lys Gln Val Gly Asp Ile Phe Ser Asn Pro Glu Thr Val Leu 260 265 270 Ala Lys Leu Ile Gln Asn Val Phe Glu Ile Lys Leu Gln Ser Phe Val 275 280 285 Lys Glu Gln Leu Glu Glu Cys Arg Lys Ser Asp Ala Glu Gln Tyr Leu 295 300 Lys Asn Leu Tyr Asp Leu Tyr Thr Arg Thr Thr Asn Leu Ser Ser Lys 315

310

Leu Met Glu Phe Asn Leu Gly Thr Asp Lys Gln Thr Phe Leu Ser Lys

```
330
                                           335
            325
Leu Ile Lys Ser Ile Phe Ile Ser Tyr Leu Glu Asn Tyr Ile Glu Val
      340 345
Glu Thr Gly Tyr Leu Lys Ser Arg Ser Ala Met Ile Leu Gln Arg Tyr
         360
                                     365
Tyr Asp Ser Lys Asn His Gln Lys Arg Ser Ile Gly Thr Gly Gly Ile
  370 375
                                 380
Gln Asp Leu Lys Glu Arg Ile Arg Gln Arg Thr Asn Leu Pro Leu Gly
               390
                              395
Pro Ser Ile Asp Thr His Gly Glu Thr Phe Leu Ser Gln Glu Val Val
            405
                            410
Val Asn Leu Leu Gln Glu Thr Lys Gln Ala Phe Glu Arg Cys His Arg
                        425
                                        430
Leu Ser Asp Pro Ser Asp Leu Pro Arg Asn Ala Phe Arg Ile Phe Thr
                            445
            440
Ile Leu Val Glu Phe Leu Cys Ile Glu His Ile Asp Tyr Ala Leu Glu
  450 455 460
Thr Gly Leu Ala Gly Ile Pro Ser Ser Asp Ser Arg Asn Ala Asn Leu
    470 475 480
Tyr Phe Leu Asp Val Val Gln Gln Ala Asn Thr Ile Phe His Leu Phe
           485 490 495
Asp Lys Gln Phe Asn Asp His Leu Met Pro Leu Ile Ser Ser Pro
                        505
Lys Leu Ser Glu Cys Leu Gln Lys Lys Glu Ile Ile Glu Gln Met
     515
                     520
                                     525
Glu Met Lys Leu Asp Thr Gly Ile Asp Arg Thr Leu Asn Cys Met Ile
                  535
Gly Gln Met Lys His Ile Leu Ala Ala Glu Gln Lys Lys Thr Asp Phe
     550
                               555
Lys Pro Glu Asp Glu Asn Asn Val Leu Ile Gln Tyr Thr Asn Ala Cys
            565
                           570
Val Lys Val Cys Ala Tyr Val Arg Lys Gln Val Glu Lys Ile Lys Asn
 580
                       585
Ser Met Asp Gly Lys Asn Val Asp Thr Val Leu Met Glu Leu Gly Val
     595 600
                                     605
Arg Phe His Arg Leu Ile Tyr Glu His Leu Gln Gln Tyr Ser Tyr Ser
                  615
                                  620
Cys Met Gly Gly Met Leu Ala Ile Cys Asp Val Ala Glu Tyr Arg Lys
                   635
               630
Cys Ala Lys Asp Phe Lys Ile Pro Met Val Leu His Leu Phe Asp Thr
            645
                            650
Leu His Ala Leu Cys Asn Leu Leu Val Val Ala Pro Asp Asn Leu Lys
                        665 670
Gln Val Cys Ser Gly Glu Gln Leu Ala Asn Leu Asp Lys Asn Ile Leu
            680 685
His Ser Phe Val Gln Leu Arg Ala Asp Tyr Arg Ser Ala Arg Leu Ala
                  695
                                  700
  690
Arg His Phe Ser
705
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     Val Asn His Ser Pro Ser Ile Ser Thr Pro Ala His Ser Ile Pro Gly
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     Ser His Ala Met
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The first state in
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     <211> 20
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     <212> PRT
Mr. ..
     <213> Homo sapiens
a Maria
     <400> 378
32 E
     Pro Glu Thr Gly Asp Pro Val Thr Leu Arg Leu Leu Asp Asp Gly Ala
      1
                                           10
1
     Gly Ala Asp Val
Brus.
Brus.
Mary Trust
=6
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      1
      His Phe Pro His
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     Leu Glu Ser Thr
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     Leu Val Thr Trp
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Ar front took house the acce.
Ar front took hour to the
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      1
Gln Ala Leu Lys
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77
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Total Street II
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1100
١. [
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Marin Shirin B
     Phe Phe Lys Trp Leu Leu Ser Cys Cys Pro Gly Ser Ser Gln Ile Ala
      1
                                             10
Ting.
     Ala Ala Ala Ser
3
                   20
Tonic cont.
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The state of
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, and
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      <213> Homo sapiens
.
.
      <400> 388
      Leu Ser Cys Cys Pro Gly Ser Ser Gln Ile Ala Ala Ser Thr Gln
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      Pro Glu Asp
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      Lys Lys Ser Gln
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Lys Met Arg Glu
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Asp Ile Asn Thr Gln Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val
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Thr Asp Ser Pro
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Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val Thr Asp Ser Pro Gly
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Arg Pro Arg Glu
            20
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Thr Ile Pro Gln
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<213> Homo sapiens
<400> 394
Val Thr Asp Ser Pro Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr
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     Ser Ser His Gly
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     Asn Arg Phe
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fr. d.
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     Met Asn Lys Leu Tyr Ile Gly Asn Leu Ser Glu Asn Ala Ala Pro Ser
17
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į
     Asp Leu Glu
122
ij
100
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ing.
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120
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1
     <213> Homo sapiens
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     Lys Ile Pro Val
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     Lys Thr Gly Tyr
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Ala Leu Ser Gly
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Lys Pro Ile Glu
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Lys Ile Glu Leu His Gly Lys Pro Ile Glu Val Glu His Ser Val Pro
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Lys Arg Gln Arg
            20
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<211> 20
<212> PRT
<213> Homo sapiens
<400> 403
Val Glu His Ser Val Pro Lys Arg Gln Arg Ile Arg Lys Leu Gln Ile
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17

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     Arg Asn Ile Pro
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     <210> 404
     <211> 20
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     <213> Homo sapiens
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     Ile Arg Lys Leu Gln Ile Arg Asn Ile Pro Pro His Leu Gln Trp Glu
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     1
     Val Leu Asp Ser
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4 12
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ı,
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١...
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123
     Ala Val Val Asn Val Thr Tyr Ser Ser Lys Asp Gln Ala Arg Gln Ala
Alin 6
     1
                                10
113
     Leu Asp Lys Leu
:
                20
100
and a
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     Asn Phe Thr Leu
                 20
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     <213> Homo sapiens
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     Asp Glu Thr Ala
                 20
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     <213> Homo sapiens
     <400> 409
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The first
     Gln Arg Gly Ser
                  20
1.3
     <210> 410
7 ......
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     <212> PRT
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ä
The Mark
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     Gly Arg Arg Gly Leu Gly Gln Arg Gly Ser Ser Arg Gln Gly Ser Pro
10
Ann day
     Gly Ser Val Ser
                  20
i nà
     <210> 411
     <211> 20
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     Ser Arg Gln Gly Ser Pro Gly Ser Val Ser Lys Gln Lys Pro Cys Asp
     1
                                           10
     Leu Pro Leu Arg
                  20
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     <213> Homo sapiens
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     Lys Gln Lys Pro Cys Asp Leu Pro Leu Arg Leu Leu Val Pro Thr Gln
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                                    10
    Ala Thr Ile Arg
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    <211> 20
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113
    <213> Homo sapiens
ij
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Ile Ile Gly Lys Glu Gly Ala Thr Ile Arg Asn Ile Thr Lys Gln Thr
1
                           10
Gln Ser Lys Ile
               20
£
25 25 E
Harry
Care
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    <213> Homo sapiens
į,
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    Asn Ile Thr Lys Gln Thr Gln Ser Lys Ile Asp Val His Arg Lys Glu
     1
     Asn Ala Gly Ala
               20
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     <211> 20
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     1
     Ile Leu Ser Thr
               20
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                        5
                                            10
      Ala Cys Lys Ser
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      <213> Homo sapiens
      <400> 418
Derit Jest grete in Bank Briefe
      Pro Glu Gly Thr Ser Ala Ala Cys Lys Ser Ile Leu Glu Ile Met His
                                            10
      Lys Glu Ala Gln
                   20
÷ ... #
      <210> 419
٠
پرد
      <211> 20
That it
      <212> PRT
      <213> Homo sapiens
ğ
114
      <400> 419
i ga
Hall Brand and
      Ile Leu Glu Ile Met His Lys Glu Ala Gln Asp Ile Lys Phe Thr Glu
      Glu Ile Pro Leu
                   20
i sk
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      catctagaaa gaagegetta agatgtggca geceetette tteaagtgge tettgteetg 180
      ttgccctggg agttctcaaa ttgctgcagc agcctccacc cagcctgagg atgacatcaa 240
      tacacagagg aagaagatc aggaaaagat gagagaagtt acagactctc ctgggcgacc 300
      ccgagagett accattecte agacttette acatggtget aacagatttg tteetaaaag 360
      taaageteta gaggeegtea aattggeaat agaageeggg tteeaceata ttgattetge 420
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                                                                             455
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<211> 161
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Tyr Phe Glu Asn Phe Leu Ala Ala Trp Arg Pro Val Lys Ala Ser Asp
            20
                                25
Gly Asp Tyr Tyr Thr Leu Ala Val Pro Met Gly Asp Val Pro Met Asp
                            40
Gly Ile Ser Val Ala Asp Ile Gly Ala Ala Val Ser Ser Ile Phe Asn
                        55
Ser Pro Glu Glu Phe Leu Gly Lys Ala Val Gly Leu Ser Ala Glu Ala
                    70
                                        75
Leu Thr Ile Gln Gln Tyr Ala Asp Val Leu Ser Lys Ala Leu Gly Lys
               8.5
                                    90
Glu Val Arg Asp Ala Lys Ile Thr Pro Glu Ala Phe Glu Lys Leu Gly
            100
                                105
Phe Pro Ala Ala Lys Glu Ile Ala Asn Met Cys Arg Phe Tyr Glu Met
                            120
Lys Pro Asp Arg Asp Val Asn Leu Thr His Gln Leu Asn Pro Lys Val
                        135
Lys Ser Phe Ser Gln Phe Ile Ser Glu Asn Gln Gly Ala Phe Lys Gly
145
                    150
                                        155
Met
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ccgatgggag atgtaccaat ggatggtatc tctgttgctg atattggagc agccgtctct 180
aqcattttta attctccaqa qqaattttta qqcaaqqccq tqqqqctcaq tqcaqaaqca 240
ctaacaatac agcaatatgc tgatgttttg tccaaggctt tggggaaaga agtccgagat 300
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            20
                                 25
Lys Ile Pro Val Ser Gly Pro Phe Leu Val Lys Thr Gly Tyr Ala Phe
                             40
Val Asp Cys Pro Asp Glu Ser Trp Ala Leu Lys Ala Ile Glu Ala Leu
                        55
Ser Gly Lys Ile Glu Leu His Gly Lys Pro Ile Glu Val Glu His Ser
                    70
                                         75
Val Pro Lys Arg Gln Arg Ile Arg Lys Leu Gln Ile Arg Asn Ile Pro
                                     90
Pro His Leu Gln Trp Glu Val Leu Asp Ser Leu Leu Val Gln Tyr Gly
                                 105
            100
                                                     110
Val Val Glu Ser Cys Glu Gln Val Asn Thr Asp Ser Glu Thr Ala Val
        115
                            120
```

** 7	~		m.1	m	0	_	-		~ 1	70.7	-	~ 1	70.7	-	70
Val	Asn 130	Val	Thr	Tyr	Ser	Ser 135	Lys	Asp	GIn	Ala	Arg 140	GIn	Ala	Leu	Asp
Lys 145	Leu	Asn	Gly	Phe	Gln 150	Leu	Glu	Asn	Phe	Thr 155	Leu	Lys	Val	Ala	Tyr 160
Ile	Pro	Asp	Glu	Thr 165	Ala	Ala	Gln	Gln	Asn 170	Pro	Leu	Gln	Gln	Pro 175	Arg
Gly	Arg	Arg	Gly 180	Leu	Gly	Gln	Arg	Gly 185	Ser	Ser	Arg	Gln	Gly 190	Ser	Pro
Gly	Ser	Val 195	Ser	Lys	Gln	Lys	Pro 200	Cys	Asp	Leu	Pro	Leu 205	Arg	Leu	Leu
Val	Pro 210	Thr	Gln	Phe	Val	Gly 215	Ala	Ile	Ile	Gly	Lys 220	Glu	Gly	Ala	Thr
Ile 225	Arg	Asn	Ile	Thr	Lys 230	Gln	Thr	Gln	Ser	Lys 235	Ile	Asp	Val	His	Arg 240
Lys	Glu	Asn	Ala	Gly 245	Ala	Ala	Glu	Lys	Ser 250	Ile	Thr	Ile	Leu	Ser 255	Thr
Pro	Glu	Gly	Thr 260	Ser	Ala	Ala	Cys	Lys 265	Ser	Ile	Leu	Glu	Ile 270	Met	His
Lys	Glu	Ala 275	Gln	Asp	Ile	Lys	Phe 280	Thr	Glu	Glu	Ile	Pro 285	Leu	Lys	Ile
Leu	Ala 290	His	Asn	Asn	Phe	Val 295	Gly	Arg	Leu	Ile	Gly 300	Lys	Glu	Gly	Arg
Asn 305	Leu	Lys	Lys	Ile	Glu 310	Gln	Asp	Thr	Asp	Thr 315	Lys	Ile	Thr	Ile	Ser 320
Pro	Leu	Gln	Glu	Leu 325	Thr	Leu	Tyr	Asn	Pro 330	Glu	Arg	Thr	Ile	Thr 335	Val
Lys	Gly	Asn	Val 340	Glu	Thr	Cys	Ala	Lys 345	Ala	Glu	Glu	Glu	Ile 350	Met	Lys
Lys	Ile	Arg 355	Glu	Ser	Tyr	Glu	Asn 360	Asp	Ile	Ala	Ser	Met 365	Asn	Leu	Gln
	370					375					380		Leu		
385			_		390					395			Ser		400
Thr	Pro	Pro	Tyr	Pro 405	Gln	Phe	Glu	Gln	Ser 410	Glu	Thr	Glu	Thr	Val 415	His
			420					425					430		Gly
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aaccgcccag agtagaagat ggattggggc acgctgcaga cgatcctggg gggtgtgaac 240
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atcctcgttg tggctgcaaa ggaggtgtgg ggagatgagc aggccgactt tgtctgcaac 360
accetgeage caggetgeaa gaaegtgtge taegateaet aetteeeeat eteecaeate 420
eggetatggg ecetgeaget gatettegtg tecageeeag egeteetagt ggeeatgeae 480
gtggcctacc ggagacatga gaagaagagg aagttcatca agggggagat aaagagtgaa 540
tttaaggaca tegaggagat caaaaceeag aaggteegea tegaaggete eetgtggtgg 600
acctacacaa gcagcatctt cttccgggtc atcttcgaag ccgccttcat gtacgtcttc 660
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Ser Thr Ser Ile Gly Lys Ile Trp Leu Thr Val Leu Phe Ile Phe Arq
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         35
Ala Asp Phe Val Cys Asn Thr Leu Gln Pro Gly Cys Lys Asn Val Cys
Tyr Asp His Tyr Phe Pro Ile Ser His Ile Arg Leu Trp Ala Leu Gln
                     70
Leu Ile Phe Val Ser Ser Pro Ala Leu Leu Val Ala Met His Val Ala
Tyr Arg Arg His Glu Lys Lys Arg Lys Phe Ile Lys Gly Glu Ile Lys
                                105
Ser Glu Phe Lys Asp Ile Glu Glu Ile Lys Thr Gln Lys Val Arg Ile
        115
Glu Gly Ser Leu Trp Trp Thr Tyr Thr Ser Ser Ile Phe Phe Arg Val
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<223> PCR primer

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Thr Val Asp Cys Phe Val Ser Arg Pro Thr Glu Lys Thr Val Phe Thr
                                185
Val Phe Met Ile Ala Val Ser Gly Ile Cys Ile Leu Leu Asn Val Thr
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Glu Leu Cys Tyr Leu Leu Ile Arg Tyr Cys Ser Gly Lys Ser Lys Lys
Pro Val
225
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<213> Artificial Sequence
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Glu Lys Ser Ile Thr Ile Leu Ser Thr Pro Glu Gly Thr Ser Ala Ala

				245					250					255	
Cys	Lys	Ser	Ile 260	Leu	Glu	Ile	Met	His 265	Lys	Glu	Ala	Gln	Asp 270	Ile	Lys
Phe	Thr	Glu 275	Glu	Ile	Pro	Leu	Lys 280	Ile	Leu	Ala	His	Asn 285	Asn	Phe	Val
Gly	Arg 290	Leu	Ile	Gly	Lys	Glu 295	Gly	Arg	Asn	Leu	Lys 300	Lys	Ile	Glu	Gln
Asp 305	Thr	Asp	Thr	Lys	Ile 310	Thr	Ile	Ser	Pro	Leu 315	Gln	Glu	Leu	Thr	Leu 320
Tyr	Asn	Pro	Glu	Arg 325	Thr	Ile	Thr	Val	Lys 330	Gly	Asn	Val	Glu	Thr 335	Cys
Ala	Lys	Ala	Glu 340	Glu	Glu	Ile	Met	Lys 345	Lys	Ile	Arg	Glu	Ser 350	Tyr	Glu
Asn	Asp	Ile 355	Ala	Ser	Met	Asn	Leu 360	Gln	Ala	His	Leu	Ile 365	Pro	Gly	Leu
Asn	Leu 370	Asn	Ala	Leu	Gly	Leu 375	Phe	Pro	Pro	Thr	Ser 380	Gly	Met	Pro	Pro
Pro 385	Thr	Ser	Gly	Pro	Pro 390	Ser	Ala	Met	Thr	Pro 395	Pro	Tyr	Pro	Gln	Phe 400
Glu	Gln	Ser	Glu	Thr 405	Glu	Thr	Val	His	Leu 410	Phe	Ile	Pro	Ala	Leu 415	Ser
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Arg	Phe	Ala 435	Gly	Ala	Ser	Ile	Lys 440	Ile	Ala	Pro	Ala	Glu 445	Ala	Pro	Asp
Ala	Lys 450	Val	Arg	Met	Val	Ile 455	Ile	Thr	Gly	Pro	Pro 460	Glu	Ala	Gln	Phe
Lys 465	Ala	Gln	Gly	Arg	Ile 470	Tyr	Gly	Lys	Ile	Lys 475	Glu	Glu	Asn	Phe	Val 480
Ser	Pro	Lys	Glu	Glu 485	Val	Lys	Leu	Glu	Ala 490	His	Ile	Arg	Val	Pro 495	Ser
Phe	Ala	Ala	Gly 500	Arg	Val	Ile	Gly	Lys 505	Gly	Gly	Lys	Thr	Val 510	Asn	Glu
Leu	Gln	Asn 515	Leu	Ser	Ser	Ala	Glu 520	Val	Val	Val	Pro	Arg 525	Asp	Gln	Thr
Pro	Asp	Glu	Asn	Asp	Gln	Val	Val	Val	Lys	Ile	Thr	Gly	His	Phe	Tyr

<213> Artificial Sequence

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ttcgtggact gcccggacga gagctgggcc ctcaaggcca tcgaggcgct ttcaggtaaa 180
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Ala Ile Ile Gly Lys Glu Gly Ala Thr Ile Arg Asn Ile Thr Lys Gln

Thr 225	210 Gln	Ser	Lys	Ile	Asp 230	215 Val	His	Arg	Lys	Glu 235	220 Asn	Ala	Gly	Ala	Ala 240
Glu	Lys	Ser	Ile	Thr 245	Ile	Leu	Ser	Thr	Pro 250	Glu	Gly	Thr	Ser	Ala 255	Ala
Cys	Lys	Ser	Ile 260	Leu	Glu	Ile	Met	His 265	Lys	Glu	Ala	Gln	Asp 270	Ile	Lys
Phe	Thr	Glu 275	Glu	Ile	Pro	Leu	Lys 280	Ile	Leu	Ala	His	Asn 285	Asn	Phe	Val
Gly	Arg 290	Leu	Ile	Gly	Lys	Glu 295	Gly	Arg	Asn	Leu	Lys 300	Lys	Ile	Glu	Gln
Asp 305	Thr	Asp	Thr	Lys	Ile 310	Thr	Ile	Ser	Pro	Leu 315	Gln	Glu	Leu	Thr	Leu 320
Tyr	Asn	Pro	Glu	Arg 325	Thr	Ile	Thr	Val	Lys 330	Gly	Asn	Val	Glu	Thr 335	Cys
Ala	Lys	Ala	Glu 340	Glu	Glu	Ile	Met	Lys 345	Lys	Ile	Arg	Glu	Ser 350	Tyr	Glu
Asn	Asp	Ile 355	Ala	Ser	Met	Asn	Leu 360	Gln	Ala	His	Leu	Ile 365	Pro	Gly	Leu
Asn	Leu 370	Asn	Ala	Leu	Gly	Leu 375	Phe	Pro	Pro	Thr	Ser 380	Gly	Met	Pro	Pro
Pro 385	Thr	Ser	Gly	Pro	Pro 390	Ser	Ala	Met	Thr	Pro 395	Pro	Tyr	Pro	Gln	Phe 400
Glu	Gln	Ser	Glu	Thr 405	Glu	Thr	Val	His	Leu 410	Phe	Ile	Pro	Ala	Leu 415	Ser
Val	Gly	Ala	Ile 420	Ile	Gly	Lys	Gln	Gly 425	Gln	His	Ile	Lys	Gln 430	Leu	Ser
Arg	Phe	Ala 435	Gly	Ala	Ser	Ile	Lys 440	Ile	Ala	Pro	Ala	Glu 445	Ala	Pro	Asp
Ala	Lys 450	Val	Arg	Met	Val	Ile 455	Ile	Thr	Gly	Pro	Pro 460	Glu	Ala	Gln	Phe
Lys 465		Gln	Gly	Arg	Ile 470		Gly	Lys	Ile	Lys 475	Glu	Glu	Asn	Phe	Val 480
Ser	Pro	Lys	Glu	Glu 485	Val	Lys	Leu	Glu	Ala 490	His	Ile	Arg	Val	Pro 495	Ser
Phe	Ala	Ala	Gly 500	Arg	Val	Ile	Gly	Lys 505	Gly	Gly	Lys	Thr	Val 510	Asn	Glu

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die Rem B
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mer mm.
Book M. B
                                             10
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     Ala Leu Ser Gly
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The Test
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                                              10
                                                                    15
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<400> 455

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Gln Arg Gly Ser
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The thirty that the
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     <213> Homo sapiens
1.3
1,3
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ž
100
     Cys Ala Lys Ala
i Pi
                  20
Arma
Trans
17
     <210> 462
ž už
     <211> 20
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     <213> Homo sapiens
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                                            10
     Ala Ser Met Asn
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                            Ile Thr Gly Pro
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                            <210> 465
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Best that the action in the control of the control 
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 Thus Thus
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```

and the second second